8/24/06

T. Gibbs 09/436060 SID 16

SCORE OVER LENGTH SEARCHES

Attached is a score over length search. This search was developed to overcome limitations in most standard search systems which favor large sequences with high scoring, but lesser overall identity over smaller sequences with higher overall identity. This search is especially useful for relatively small nucleic acid or polypeptide target sequences (antisense, fragments, probes, primers, RNAi, epitopes, haptens, etc.) claimed functionally via a form of hybridization and/or identity language and having defined upper and lower polynucleotide and or polypeptide length limits.

The score over length search is performed by first running the query sequence using examiner-specified identity and polynucleotide or protein length limit parameters, and saving 65,000 hits and 0 alignments from each desired database. The resulting output is reformatted using a Microsoft Word macro and is imported into Excel. The summary table data are then sorted by the ratio of score of each hit sequence divided by its length and the accession numbers for all hits below the examiner's desired score over length parameters are deleted. The remaining accession numbers are used to pull the corresponding sequences from the databases into subdatabases enriched for good hits and the query sequence is re-run against these subdatabases to yield the final results.

The score over length cutoff for this search is 75.%

Examiner Please Note: This cover sheet should be included when submitting results to be scanned.

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STIC-Biotech/ChemLib

161786

From:

Gibbs, Terra

Sent:

Monday, August 08, 2005 12:05 PM

To: Subject:

STIC-Biotech/ChemLib sequence search request...

I have another request for a score over length search:

I need a length limited nucleotide sequence search of SEQ ID NO:16 in USSN 09/436,060, where the returns are rank ordered based on the score over length/ratio as we've discussed. I need the lengths limited to hits between 8 and 80 nucleotides, and I'll take as many hits as you can import into excel (64,000?), and alignments for anything above .75 on the above ratio. Hope this is clear, please call me if it's not. I also need the interference databases searched.

Terra Cotta Gibbs, Ph.D. Art Unit 1635 Remsen Building 2D10 Mailbox 2C18 571-272-0758

STAFF USE ONLY

Searcher:
Searcher Phone: 2Date Searcher Picked up:
Date Completed:
Searcher Prep/Rev. Time:
Online Time:

Type of Search

NA#:____ AA#:___
Interference:___ SPDI:__
S/L:___ Oligomer:___
Encode/Transl:___ Text:__
Structure#:____ Text:__
Inventor:___ Litigation:___

QUESTEL/ORBIT:______
LEXIS/NEXIS:_____
SEQUENCE SYSTEM:_____

WWW/Internet:_____ Other(Specify):_____

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ACCESSION: AR3 06472 ACCESSION: AR3 73 06 ACCESSION: AX4 65471 ACCESSION: BD023701 ACCESSION: BA955 ACCESSION: A84 596 ACCESSION: AR0 79889 ACCESSION: AR0 79892 ACCESSION: ARO 79899	ACCESSION: BD058136 ACCESSION: BD058137 ACCESSION: BD058137 ACCESSION: AR44592 ACCESSION: AR279618 ACCESSION: AR370168 ACCESSION: AR370168 ACCESSION: AR4594 ACCESSION: AR4594 ACCESSION: AR479891 ACCESSION: AR679891 ACCESSION: AR67663850 ACCESSION: AR676653 ACCESSION: AR676653 ACCESSION: AR676653 ACCESSION: AR676653 ACCESSION: AR676653	ACCESSION: BD176174 ACCESSION: A494987 ACCESSION: A494988 ACCESSION: A8016055 ACCESSION: AR016061 ACCESSION: AR028785 ACCESSION: AR028786 ACCESSION: AR028786 ACCESSION: AR028716 ACCESSION: AR059216 ACCESSION: AR05527 ACCESSION: AR075527 ACCESSION: AR075527 ACCESSION: AR075533	ACCESSION: BDI 7616 ACCESSION: BDI 76171 ACCESSION: BDI 76171 ACCESSION: BDI 6107 ACCESSION: BD 6508 ACCESSION: BD 6468 ACCESSION: AR243518 ACCESSION: AR243519 ACCESSION: AR3 06473 ACCESSION: AR3 06473 ACCESSION: AR3 0169 ACCESSION: AR3 90720 ACCESSION: AR3 90721 ACCESSION: AR3 90721 ACCESSION: AR3 90721 ACCESSION: AR3 90731
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	Local Similarity 100.0%; Pred. No. 1.4; ss 54; Conservative 0; Mismatches 0; Indels 1 GGGTTGCGGAGGGGGGGGGGGGGGGGGGGGGGGGGGGGG	RESULT 2 AR154150 LOCUS DEFINITION Sequence 15 from patent US 6238867. ACCESSION AR154150 VERSION AR154150. KEYWORDS SOURCE THE SECURAL SEQUENCE OF THE SECURAL SEQUENCE OF THE SECURAL SECURA	NISM NCE ORS E NAL NAL ES	Query Match 12.0%; Score 54; DB 1; Length 66; Best Local Similarity 100.0%; Pred. No. 1.5; Astches 54; Conservative 0; Mismatches 0; Indels 0; Gaps 0; Qy 1 GGGTTGCGGAGGGTGGGCCTGGGAGGGTGGTGGTGGCCATTTTTTGTCTAACCCTA 54		AUTHORS Keith, W.N. TITLE Promoter region of mouse and human telomerase RNA component genes JOURNAL CANCER RESEARCH CAMPAIGN TECHNOLOGY LTD COMMENT OS AFTIFICIAL Sequence PN JP 200250969-A/82 PD 02-APR-2002 PP 02-APR-1099 JP 200529424 PR 29-JAN-1999 GB 9801902.9 PI WILLIAM NICOL KEITH PC 29-JAN-1999 GB 9801902.9 PI WILLIAM NICOL KEITH PC 212N15/09, A61K31/7105, A61K31/711, A61K35/76, A61K38/00, A61K48/00, PC A61F35/00, C12N1/15, C12N1/19, C12N1/21, C12N5/10, C12P21/02 PC C12Q1/68//C12N9/12,
ACCESSION: AX423145 ACCESSION: AX735544 ACCESSION: AX735544 ACCESSION: AX781412 ACCESSION: AX786941 ACCESSION: BD071035 ACCESSION: BD071046 ACCESSION: BD071076 ACCESSION: BD071076	ACCESSION: BD071085 ACCESSION: BD071088 ACCESSION: BD071089 ACCESSION: BD533144 ACCESSION: BD253907 ACCESSION: BD253907 ACCESSION: BD266444	ACCESSION: BD258356 ACCESSION: BD258356 ACCESSION: BD273168 ACCESSION: AR224417 ACCESSION: AX214417 ACCESSION: AX214566 ACCESSION: AX214616 ACCESSION: AX214617 ACCESSION: AX214617 ACCESSION: AX214617	ACCESSION: AX226635 ACCESSION: AX226635 ACCESSION: AX224613 ACCESSION: AX324614 ACCESSION: AX422502 ACCESSION: AX422503 ACCESSION: AX531207 ACCESSION: AX531207 ACCESSION: AX54525 ACCESSION: AX54525 ACCESSION: AX54525	ACCESSION: AX688083 ACCESSION: BD104967 ACCESSION: BD25841 ACCESSION: AX019590 ACCESSION: AX063826 ACCESSION: BD225839 ACCESSION: AX019588 ACCESSION: AX19588	ACCESSION: ARLE 41.50 ACCESSION: AX019629 ACCESSION: AX019629 ACCESSION: BD225882 ACCESSION: BAX019631 ACCESSION: AX019631 ACCESSION: AX041868 ACCESSION: AX940863 ACCESSION: AX3840863 ACCESSION: AX384137 ACCESSION: AX327120	linear PAT 08-AUG-2001
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FEATURES

PAT 17-JUL-2003

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bp DNA linear PAT 17-JUL-200 human telomerase RNA component genes.
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Patent: UP 2002209699-A 85 02-APR-2002;
CANCER RESEARCH CAMPAIGN TECHNOLOGY LTD
OS Artificial Sequence
                                                                                             Promoter regions of the mouse and human telomerase rna component
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KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH (GB)
Location/Qualifiers
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KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH
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Best Local Similarity 94.1%; Pred. No. 4.2;
Matches 48; Conservative 0; Mismatches 3;
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
/note="Mutant construct"

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Promoter region of mouse and human telomerase RNA component genes.
BD225880
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Promoter region of mouse and human telomerase RNA component genes Patent: 19 2002509699-A 83 02-APR-2002;
CANCER RESEARCH CAMPAIGN TECHNOLOGY LTD
OS Artificial Sequence
PN JP 2002509699-A/83
PD 02-APR-2002
PP 29-JAN-1999 JP 2000529424
PR 29-JAN-1999 GB 9801902.9
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PC A61P35/00,
'C1201/68//C12N9/12,
PC (A61K35/76,A61K31:522),C12N15/00,A61K37/02,C12N5/00 CC
Description of Artificial Sequence: Mutant construct FH Key
Location/Qualifiers
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Description of Artificial Sequence: Mutant construct FH
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                                                                                                                                                                                                        Score 46.2; DB 1;
Pred. No. 4.2;
0; Mismatches 3;
                                                                                                                                 /organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
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/organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   synthetic construct
synthetic construct
other sequences; artificial sequences.
1 (bases 1 to 51)
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larity 94.1%; Pred. No. 4.2;
Conservative 0; Mismatches
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Sequence 82 from Patent W09938964.
AX019628 GI:10043542
                                                         . 51
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BD225880.1 GI:33035650
                                                                                                                                                                                                          th 10.2%;
Similarity 94.1%;
48; Conservative
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                                                         source
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Best Local Similarity
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Matches 48; Conserv
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PAT 07-SEP-2000
                                                                                                          Artificial Sequence

Artificial Sequence

N JP 2002509699-A/84

DO 02-APK-2002

PF 29-JAN-1999 UP 2000529424

PR 29-JAN-1999 GB 9801902.9

PI WILLIAM NICOL KEITH

PC
C12N15/09,A61K31/7105,A61K31/711,A61K35/76,A61K38/00,A61K45/00, PC
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Promoter region of mouse and human telomerase RNA component genes.
BD225883
                                       Promoter region of mouse and human telomerase RNA component genes Patent: JP 2002509699-A 84 02-APR-2002; CANCER RESEARCH CAMPAIGN TECHNOLOGY LTD OS Artificial Sequence PN JP 2002509699-A/84

PD 02-APR-2002
                                                                                                                                                                                                                                                                                                                                      PC A61P95/00,C12N1/15,C12N1/19,C12N1/21,C12N5/10,C12P21/02 PC C12Q1/68/C12N9/12, PC (A6KX3/76,A61K31:522),C12N15/00,A61K37/02,C12N5/00 CC Description of Artificial Sequence: Mutant construct FH Key Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Promoter regions of the mouse and human telomerase rna component
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KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH (GB)
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 GGGTTGCGGAGGGCCTGGGAGGGGTGGTGGCCATTTTTGTCTAACC
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Pred. No. 9.4;
0; Mismatches 6; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /organism='Artificial Sequence' Location/Qualifiers
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/organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="texon:32630"
/note="Mutant construct"
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    .51
    /organism="synthetic col/mol_type="genomic DNA"
    /db_xref="taxon:32630"

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88.2%;
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BD225883
LOCUS
DEFINITION
ACCESSION
VERSION
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               AUTHORS
TITLE
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AX019630
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TITLE
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PN JP 40022

PD 02-APR-2002

PF 29-ANN-1998 GB 9801902.9

PI WILLIAM NICOL KEITH

PC C12N15/09,A61K31/7105,A61K31/711,A61K35/76,A61K38/00,A61K45/00, PC A61K48/00,

PC A61K35/00,C12N1/15,C12N1/19,C12N1/21,C12N5/10,C12P21/02 PC A61K35/76,A61K31:522),C12N1/19,C12N1/21,C12N5/10,C12P21/02 PC C12N1/68/C12N9/12.

PC A61K35/76,A61K31:522),C12N15/00,A61K37/02,C12N5/00 CC PC CA61K35/76,A61K31:522),C12N15/00,A61K37/02,C12N5/00 CC PC CA61K35/76,A61K31:522),C12N15/00,A61K37/02,C12N5/00 CC PC A61K35/76,A61K31:522) C12N15/00,A61K37/02,C12N5/00 CC PC A61K35/76,A61K31/02 A61K35/76 A61
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Promoter region of mouse and human telomerase RNA component genes.
BD225881
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KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              9.5%; Score 43; DB 1; Length 51;
larity 90.2%; Pred. No. 7.2;
Conservative 0; Mismatches 5; Indels
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/organism="synthetic construct"
/mol_type="unassigned DNA"
/db xref="taxon:32630"
/noCe="Mutant construct"
                                                                                                                                                                                                                                                                                                                                                                                                                    1..51
/organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
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synthetic construct
synthetic construct
other sequences; artificial sequences.
1 (bases 1 to 51)
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Sequence 85 from Patent WO9938964.
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synthetic construct
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Matches 46; Conserv
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Promoter region of mouse and human BD225837
                                                      BD225837.1 GI:33035607
JP 2002509699-A/40.
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                                                                                     synthetic construct
synthetic construct
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Matches 36; Conserv
                                                                                                                                              Keith, W.N.
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TITLE
JOURNAL
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VERSION
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AX019586
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                                                                                    Promoter region of mouse and human telomerase RNA component genes
Patent: JP 2002509699-A 86 02-APR-2002;
CANCER RESEARCH CAMPAIGN TECHNOLOGY LTD
OS Artificial Sequence
PN JP 2002509699-A/86
PD 02-APR-2002
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PC (A61K35/76,A61K31:522),C12N15/00,A61K37/02,C12N5/00 CC Description of Artificial Sequence: Mutant construct FH Key Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH
Location/Qualifiers
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/organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
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    .51
    ^coganism="synthetic construct"
|mol_type="unassigned DNA"
|db_xref="taxon:32630"

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                                          other sequences; artificial sequences.
1 (bases 1 to 51)
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Pred. No. 16;
0; Mismatches
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34.3%; Pred. No. 16;
ive 0; Mismatches
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Sequence 86 from Patent W09938964.
AX019632
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                                                                                                                                                                           29-JAN-1999 JP 2000529424
29-JAN-1998 GB 9801902.9
WILLIAM NICOL KEITH
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1 Similarity 84.3%;
43; Conservative (
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Conservative
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JP 2002509699-A/86.
synthetic construct
synthetic construct
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Best Local Similarity
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                                                                       Keith, W.N.
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Promoter region of mouse and human telomerase RNA component genes Patent: JP 200259659-A 40 02-APR-2002; CANCER RESEARCH CAMPAIGN TECHNOLOGY LTD
CANCER RESEARCH CAMPAIGN TECHNOLOGY LTD
CANCER RESEARCH CAMPAIGN TECHNOLOGY LTD
CO-APR-2002
PD 02-APR-2002
PP 29-JAN-1999 JP 2000529424
PR 29-JAN-1998 GB 9801902.9
PI WILLIAM NICOL KEITH
PC C12N15/09,A61K31/7105,A61K31/711,A61K35/76,A61K38/00,A61K45/00, PC
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DNA linear PAT 17-JUL-200 telomerase RNA component genes.
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PC A61P35/00,C12N1/15,C12N1/19,C12N1/21,C12NS/10,C12P21/02,

C12Q1/68/C12N9/12,

PC (A61K35/76,A61K31:522),C12N15/00,A61K37/02,C12NS/00 CC

Description of Artificial Sequence:Oligonucleotide FH Key

Location/Qualifiers
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Location/Qualifiers
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KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH
Location/Qualifiers
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17;
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/organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
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/noTe="Oligonucleotide"
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(bases 1 to 38)
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synthetic construct
other sequences; artificial sequences.
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Sequence 40 from Patent WO9938964.
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A61K48/00,
PC A61P35/00,C12N1/15,C12N1/19,C12N1/21,C12N5/10,C12P21/02 PC,C12Q1/68//C12N9/12,
C12Q1/68//C12N9/12,
PC (A61K35/76,A61K31:522),C12N15/00,A61K37/02,C12N5/00 CC
Description of Artificial Sequence:Oligonucleotide FH Key
Location/Qualifiers
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KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH (GB)
Location/Qualifiers
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KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH (GB)
Location/Qualifiers
1. .38
                                                                                                                             /organism='Artificial Sequence'
                                                                                                                                                                                                                                                                   Length 38;
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/noTe="Oligonucleotide"
                                                                                                                                                                        1. 38
/organism="synthetic construct"
/mol type="genomic DNA"
/db_kref="taxon:32630"
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6.9%; Score 31.2; D
Best Local Similarity 91.7%; Pred. No. 36;
Matches 33; Conservative 0; Mismatches
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Sequence 42 from Patent W09938964.
AX019588
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Sequence 41 from Patent WO9938964.
AX019587
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AX019588
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PN JP 2002509699-A/41
PD 02-APR-2002
PP 29-JAN-1999 GB 9801902.9
PR 29-JAN-1999 GB 9801902.9
PI WILLIAM NICOL KEITH
PC
C12N15/09, A61K31/7105, A61K31/711, A61K35/76, A61K38/00, A61K45/00, PC
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Promoter region of mouse and human telomerase RNA component genes.
BD225838
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CANCER RESEARCH CAMPAIGN TECHNOLOGY LTD
S Artificial Sequence
N JP 2003509699-A/42
PD 02-APR-2002
PP 29-JAN-1999 JP 2000529424
PF 29-JAN-1999 GB 9801902.9
PI WILLIAM NICOL KEITH
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                                                                                                                                                                                                                                                                                                Promoter region of mouse and human telomerase RNA component genes Patent: JP 200250959-A 41 02-APR-2002;
CANCER RESEARCH CAMPAIGN TECHNOLOGY LTD
OS Artificial Sequence
PN JP 2002509699-A/41
PD 02-APR-2002
PR 29-JAN-1998 JP 2000529424
PR 29-JAN-1998 GB 9801902.9
PI WILLIAM NICOL KEITH
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PC A61P35/00,C12N1/15,C12N1/19,C12N1/21,C12N5/10,C12P21/02 PC (A61P35/00,C12N1/15,C12N1/19,C12N1/21,C12N5/10,C12P21/02 PC (A61K35/76,A61K31:522),C12N15/00,A61K37/02,C12N5/00 CC Description of Artificial Sequence:Oligonucleotide FH Key Location/Qualifiers
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Location/Qualifiers
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    .38
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    /db_xref="taxon:32630"

GGGTTGCGGAGGGTGGGCCTGGGAGGGGTGGTGGCC
                                                                                                                                                                                                                                          other sequences, artificial sequences.
1 (bases 1 to 38)
Keith, W.N.
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synthetic construct
other sequences; artificial sequences.
1 (bases 1 to 38)
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1 Similarity 91.7%; Pred. No. 36;
33; Conservative 0; Mismatches
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BD225839
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BD225838
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PAT 07-SEP-2000

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bu071082 31 bp RNA linear PAT 27-AUG-2002
Modulation of mammalian telomerase by peptide nucleic acids.
BD071082
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                                                                                                                                                                                                                     Unidentified
JP 2001517929-A/48
09-OCT-2001
09-APR-1997 JP 1997536487
09-APR-1996 US 08/630019
JERRY W SHAY, WOODRING E WRIGHT, MIECZYSLAW A PIATYSZEK, DAVID
                                                                                                                                       1 (bases 1 to 31)
Shay, J.W., Wright, W.E., Piatyszek, M.A., Corey, D. and Norton, J.C.
Modulation of mammalian telomerase by peptide nucleic acids
Patent: JP 2001517929-A 48 09-OCT-2001;
GERON CORP
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Unclassified.
1 (bases 1 to 30)
1 (laponteau, B., Feng, J., Funk, W. and Andrews, W.H. Assays for the DNA component of human telomerase Patent: US 5776679-A 22 07-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1; Length 31; 30;
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Sequence 22 from patent US 5776679.
AR016054
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    .31
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Strandedness: Single;
Topology: Linear;
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                                                            BD071082.1 GI:22616685
JP 2001517929-A/48.
unidentified
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JAMES C NORTON
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Weinrich, S.L., Atkinson, E.M. III, Lichtsteiner, S.P., Vasserot, A.P.
and Pruzan, R.A.
Purified telomerase
Patent: US 6517834-A 1 11-FEB-2003;
Location/Qualifiers
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Weinrich, S.L., Atkinson, E.M. III, Lichtsteiner, S.P., Vasserot, A.P.
Meinrich, S.L., Atkinson, E.M. III, Lichtsteiner, S.P., Vasserot, A.P.
Methods for purifying telomerase
Methods for purifying telomerase
Patent: US 6545133-A 1 08-APR-2003;
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                                                                                 DB 1;
 /organism="synthetic construct"
/mol type="unassigned DNA"
/db_xref="taxon:32630"
/nore="Oligonucleotide"
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Sequence 1 from patent US 6545133.
AR305067
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/organism="unknown"
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1 Similarity 91.7%;
33; Conservative
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AR305067/c
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1 (bases 1 to 30)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H. Mammalian telomerase
Patent: US 6258535-A 22 10-JUL-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 30;
          Length 30;
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Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Wamman, I an telomerase
Patent: US 5958680-A 23 28-SEP-1999;
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                                                                137 CCTGCCGCCTTCCACCGTTCATTCTAGAGC 166
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llarity 100.0%; Pred. No. 34;
Conservative 0; Mismatches
        Query Match 6.7%; Score 30; DB Best Local Similarity 100.0%; Pred. No. 34; Matches 30; Conservative 0; Mismatches
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Sequence 23 from patent US 5958680.
AR075526.1 GI:10002274
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Sequence 22 from patent US 6258535.
AR161924
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Mammalian telomerase.
BD176165
BD176165.1 GI:29121871
JP 2002272489-A/24.
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                                                                                                                                                                                                                                                               Unclassified.
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Kim,N.Woo., Wu.F., Kealey,J.T., Pruzan,R. and Weinrich,S.L.
Methods for detecting the RNA component of telomerase
Patent: US 5846723-A 8 08-DEC-1998;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unknown...
Unclassified.
Unclassified.
I (bases 1 to 30)
Kim,N.Woo., Wu,F., Kealey,J.T., Pruzan,R. and Weinrich,S.L.
Kim,N.Woo., Wu,F., Kealey,J.T., Pruzan,R. and Weinrich,S.L.
Methods for decting the RNA component of telomerase
Patent: US 5846723-A 5 08-DEC-1998;
Location/Qualifiers
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1 (bases 1 to 30)
11leponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Mammalian telomerase
Patent: US 5837857-A 22 17-NOV-1998;
Locaion/Qualifiers
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100.0%; Pred. No. 34;
iive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                         290 CTGCCACCGCGAAGAGTTGGGCTCTGTCAG 319
           AR059215
Sequence 22 from patent US 5837857.
AR059215.1 GI:5984792
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/organism="unknown"
/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
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Sequence 5 from patent US 5846723.
AR063829.1 GI:5993137
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Sequence 8 from patent US 5846723.
AR063832.1 GI:5993140
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Best Local Similarity 100.09
Matches 30, Conservative
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Best Local Similarity 100.
Matches 30; Conservative
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AR063829/c
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AR063832/c
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AR059215/c
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PAT 10-APR-2003

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1 (bassel to 30)
Weinrich,S.L., Atkinson,E.M. III, Lichtsteiner,S.P., Vasserot,A.P., and Pruzan,R.A.
Purified telomerasa.
Purified telomerasa.
Purified telomerasa.
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Weinrich, S.L., Atkinson, E.M. III, Lichtsteiner, S.P., Vasserot, A.P.
and Pruzan, A.A.
Purified telomerase
Patent: US 6517834-A 5 11-FEB-2003;
Location/Qualifiers
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Weinrich, S.L., Atkinson, E.M. III, Lichtsteiner, S.P., Vasserot, A.P.
and Pruzan, R.A.
Methods for purifying telomerase
Patent: US 6545133-A 4 08-APR-2003;
Location/Qualifiers
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100.0%; Pred. No. ...
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Sequence 5 from patent US 6517834.
AR279620 GI:29714514
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/wol_type="genomic DNA"
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/organism="unknown"
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Best Local Similarity 100.*
Matches 30; Conservative
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Matches 30; Conservative
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AR279620/c
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AR305070/c
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PN JP 2002272489-A/24
PN JP 2002272489-A/24
PD 24-SEP-2002 JP 2002061125
PF 06-MAR-2002 JP 2002061125
PR 07-JUJ-1994 US 08/372102,27-OCT-1994 US 08/330123 PR 07-JUJ-1995 US 08/472802,07-JUN-1995 US 08/482115 PI BRYANT VILLEPONTEAL,JUNIA FENO,WALTER FUNK,WILLIAM H ANDREMS PC C12N15/09,C12N9/99,C12Q1/68,G01N33/53,G01N33/56,C12N15/00 CC Strandedness: Single, Linear;
CC Mammalian telomerase
CC Mammalian telomerase
FH Sey Location/Qualifiers
FT source /organism='Unidentified'.
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Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Mammalian telomerase
Patent: JP 2002272489-A 24 24-SEP-2002;
GERON CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             6.7%; Score 30; DB 1; Length 30; larity 100.0%; Pred. No. 34; Conservative 0; Mismatches 0; Indels
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Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H. Mammalian telomerase
Patent: US 5583016-A 22 10-DEC-1996;
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/organism='Unidentified'
Location/Qualifiers
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Sequence 22 from patent US 5583016.
131769
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/mol_type="unassigned DNA"
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Sequence 4 from patent US 6517834.
AR279619
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Matches 30; Conserv
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PAT 12-JUN-2003

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RESULT 30 AR279619/c

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PAT 16-JUL-2002

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TITLE JOURNAL FEATURES

AR305071/c

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BD023701 30 bp DNA linear PAT 27-AUG-2002
Method for detecting and inhibiting RNA component of telomerase.
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unclassified.
1 (bases 1 to 30)
Extent JP 2001507229-A 5 05-JUN-2001;
Patent JP 2001507229-A 5 05-JUN-2001;
PP 19-DEC-1997 JP 1998529003
PP 19-DEC-1996 US 08/770565 PI
NAM WOO KIM, FRED WU, JAMES T KEALEY, RONALD PRUZAN, SCOTT L PI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Letsinger, R.L. and Garimella, V.
Method of detection by enhancement of silver staining
Patent: WO 0204681-A 2 17-3AN-2002;
NORTHWEFERN UNIVERSITY (US)
Location/Qualifiers
1 (bases 1 to 30)
Letsinger.R.L. and Garimella,V.
Method of detection by enhancement of silver staining
Patent: 10 5602669-A 2 05-AUG-2003;
Location/Qualifiers
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                                                                                                                                                Length 30;
                                                                                                                                                                                                                                                                                                                                linear
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Best Local Similarity 100.0%; Pred. No. 34;
Matches 30; Conservative 0; Mismatches 0
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synthetic construct
other sequences; artificial sequences.
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Sequence 2 from Patent WO0204681.
AX465471
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BD023701.1 GI:22564924
JP 2001507229-A/5.
unidentified
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BD023701/c
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DEFINITION
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AX465471/c
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ORGANISM
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Weinrich, S.L., Atkinson, E.M. III, Lichtsteiner, S.P., Vasserot, A.P.
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Unclassified.
Unclassified.
1 (bases 1 to 30)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Mammalian telomerase 2 15-APR-2003;
Patent: US 6548298-A 22 15-APR-2003;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1; Length 30;
34;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              linear
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34;
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                                                                                                                                    DNA
                  137 CCTGCCGCCTTCCACCGTTCATTCTAGAGC 166
                                                                                                                                                                                                                                                                                                                                                                                                                                6.7%; Score 30; DB 100.0%; Pred. No. 34; ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 6.7%; Score 30; DB Best Local Similarity 100.0%; Pred. No. 34; Matches 30; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      30 bp
Sequence 22 from patent US 6548298.
AR306472
AR306472.1 GI:31696311
                                                                                                                                                                                                                                                                                              and Pruzan, R.A.
Methods for purifying telomerase
Patent: US 654513-A 5 08-APR-2003;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CCTGCCGCCTTCCACCGTTCATTCTAGAGC 1
                                                                                                                             Sequence 5 from patent US 6545133.
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Sequence 2 from patent US 6602669.
AR373060
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/organism="unknown"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1. .30
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                                                                                                                                                                                AR305071.1 GI:31694378
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Best Local Similarity 100.0
Matches 30; Conservative
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Unclassified.
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Unclassified.
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AR306472/c
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RESULT 35 AR373060/c

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PAT 21-JAN-2000

linear

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Unclassified.

1 (bases I to 30)
Weinrich, S. L., Akkinson, E.M. III, Lichtsteiner, S.P., Vasserot, A.P., Pruzan, R.A. and Kealey, J.T.
Pruzan, R.A. and Kealey, J.T.
Purified telhomerase
Parent: US 5968506-A 2 19-OCT-1999;
Location/Qualifiers
                                                                                                                                                                                                                             DB 1; Length 30; 40;
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Best Local Similarity 100.0%; Pred. No. 40;
Matches 29; Conservative 0; Mismatches C
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 Atkinson, E.M. and Kealey, J.T.
PURIFIED TELCMERASE
Patent: WO 9845450-A 5 15-OCT-1998;
GERON CORP (US)
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    /organism="unidentified"
/mol_type="unassigned DNA"
/db_xref="taxon:32644"

                                                                                                                                                                                                                                                                                                  167 AAACAAAAATGTCAGCTGCTGGCCCGTT 195
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Patent: WO 9845450-A 6 15-OCT-1998;
GERON CORP (US)
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                                                                                                                                                                             /note="N = BIOTINYLATED G"
/mod_base=OTHER
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/mod_base=OTHER
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Squence 2 from patent US 5968506.
AR079889.1 GI:10006642
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Sequence 6 from Patent WO9845450.
A84596.1 GI:6733512
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Atkinson, E.M. and Kealey, J.T.
PURIFIED TELOMERASE
                                                                         Location/Qualifiers
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/organism="unknown"
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unclassified.
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AR079889/c
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A84596/c
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PD JP 2001507229-A/8

PD 05-JUN-2001;

PF 19-DEC-1997 JP 1998529003

PR 20-DEC-1996 US 08/770564,20-DEC-1996 US 08/770565 PI
NAM WOO KIM, FRED WU, JAMES T KEALEY, RONALD PRUZAN, SCOTT L. PI
WEINRICH

PC C12N19/10, A61K9/08, A61K31/7105, A61K45/00, A61K48/00, A61P35/00, PC C12N15/10, PC C12N19/12, C12Q1/68, C12N15/00, C12N19/12, C12Q1/68, C12N15/00, C12N19/12, Linear;

CC 70pology: Linear;
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Kim_N.W., Wu.F., Kealey,J.T., Fruzan,R. and Weinrich,S.L.
Method for detecting and inhibiting RNA component of telomerase
Patent: JP 2001507229-A 8 05-JUN-2001;
GERON CORP
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                                                                                                                                                                             6.7%; Score 30; DB 100.0%; Pred. No. 34; tive 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /organism="unidentified"
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Sequence 5 from Patent WO9845450.
A84595
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                                                                        Location/Qualifiers
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JP 2001507229-A/8.
unidentified
unidentified
Strandedness: Single;
CC Topology: Linear;
CC /note= 'oligo 16'
FH Key
                     Topology: Linear;
/note= 'oligo 16'
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Best Local Similarity 100.0
Matches 30; Conservative
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unclassified.
1 (bases 1 to 30)
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A84595/c
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BD023704/c
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PAT 31-AUG-2000

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Eukaryots; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;

Spermatophyta; Magnoliophyta; Liliopaida; Poales; Poaceae; PACCAD
clada; Panicoideae; Andropogoneae; Zea.

I (bases 1 to 30);

St Weinrich,S.L., Iii,E.M.A., Lichtsteiner,S.P., Vasserot,A.P.,

Purified telomerase

AL Patent: JP 2001509681-A 6 24-JUL-2001;

GERON CORP

PN JP 2001509681-A/6

PN JP 2001509681-A/6

PD 24-JUL-2001

PF 04-APR-1997 JP 1998542718

PI SCOTT L WEINRICH,EDWARD M ATKINSON III,SERGE P LICHTSTEINER,

PI ALAIN P VASSEROT,RONALD A PRUZAN,JAMES T KEALEY PC

CILNIS/S4,CI2N9/12,CO7KI6/40,CI2Q1/68,CO7KI4/47 CC Strandedness:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PAT 27-AUG-2002
                                                                              Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Viridiplantae; Streptophyta; Endracophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; PACCAD clade; Panicoideae; Andropogoneae; Zea.
1 (bases 1 to 30)
Weinrich, S.L., Iii, E.M.A., Lichtsteiner, S.P., Vasserot, A.P., Pruzan, R.A., and Kealey, J.T.
                                                                                                                                                                                                                              PN 24-JUL-2001
PD 24-JUL-2001
PF 04-APR-1997 JP 1998542718
PF 3COTT L WEINRICH, EDWARD M ATKINSON III, SERGE P LICHTSTEINER, PI ALAIN P VASSEROT, RONALD A PRUZAN, JAMES T KEALEY PC
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/mod_base= OTHER
/note= 'N = biotinylated G'
/note= 'oligonucleotide 14'
*~v
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA
                                                                                                                                                                                  Purified telomerase
Patent: JP 2001509681-A 5 24-JUL-2001;
GERON CORP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA"
                                                                                                                                                                                                                                                                                                                                                     Topology: Linear;
/mod_base= OTHER
/note= 'N = biotinylated G'
/note= 'oligonucleotide 13'
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
               BD058136.1 GI:22603742
JP 2001509681-A/5.
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JP 2001509681-A/6.
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CC /moc
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FH Key
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CC Top
CC /mo
CC /no
CC /no
FH Key
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BD058137/c
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Weinrich, S.L., Atkinson, E.M. III, Lichtsteiner, S.P., Vasserot, A.P., Pruzan, R.A. and Kealey, J.T.
Pruzin, R.A. and Kealey, J.T.
Purified telomeras 5 19-0cr-1999;
Patent: Location/Qualifiers
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Weinrich, S.L., Akkinson, E.M. III, Lichtsteiner, S.P., Vasserot, A.P., Pruzan, R.A. and Kealey, J.T.
Purified telomerase
Purified telomerase
Patent: Location/Qualifiers
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/organism="unknown"
/wol_type="unassigned DNA"
 /mol_type="unassigned DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                167 AAACAAAAATGTCAGCTGCTGGCCCGTT 195
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /mol_type="unassigned DNA"
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                                                                                                                                                                                                                Sequence 5 from patent US 5968506.
                                                                                                     43 TGTCTAACCCTAACTGAGAAGGGCGTAGG 71
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 6 from patent US 5968506.
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/organism="unknown"
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Purified telomerase.
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Best Local Similarity 100.0
Matches 29; Conservative
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BD058136/c
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AR079893/c
LOCUS
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AR079892/c
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PAT 10-APR-2003
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1 (bases 1 to 30)

Wennich, S.D., Atkinson, E.M. III, Lichtsteiner, S.P., Vasserot, A.P.
Wennich, S.L.,
and Pruzan, R.A.
Methods for purifying telomerase
Patent: US 6545133-A 3 08-APR-2003;
Location/Qualifiers

1. .30
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1 (Dases 1 to 30)
Weinrich, S.L., Atkinson, E.M. III, Lichtsteiner, S.P., Vasserot, A.P. and Pruzan, R.A.
Purified telomerase
Purified telomerase
Patent: US 6517834-A 3 11-FEB-2003;
Location/Qualifiers
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Strandedness:
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PI ALAIN P VASSEROT, RONALD A PRUZAN, JAMES T KEALEY PC C12N15/54, C12N9/12, C07K16/40, C12Q1/68, C07K14/47 CC Stris Single;
CC /mod_base= OTHER
CC /mode base= OTHER
FF KEY
FF Modified base 1
FT modified base 31.
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6.3%; Score 28.4; DB 1; Length 30;
Best Local Similarity 96.7%; Pred. No. 45;
Matches 29; Conservative 0; Mismatches 1; Indels
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Best Local Similarity 100.0%; Pred. No. 42;
Matches 29; Conservative 0; Mismatches
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Sequence 3 from patent US 6545133.
AR305069
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Sequence 3 from patent US 6517834.
AR279618 GI:29714512
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/organism="unknown"
/mol_type="genomic DNA"
                                                                                                                                                                                                                                     /organism="Zea mays"
/mol_type="genomic DNA"
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AR305069/c
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AR279618/c
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Spermatophyta, Magnoliophyta, Liliopsida, Poales, Poaceae, PACCAD
clade, Panicoideae, Andropogoneae, Zea.
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24-JUL-2011
04-APR-1997 JP 1998542718
SCOTT L WEINRICH, EDWARD M ATKINSON III, SERGE P LICHTSTEINER,
                                                                                                                                          Gaps
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Weinrich, S.L., Iii, E.M.A., Lichtsteiner, S.P., Vasserot, A.P., Pruzan, R.A. and Kealey, J.T.
Purified telomerase
Patent: JP 2001509681-A 2 24-JUL-2001;
GERON CORP
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100.0%; Pred. No. 42;
tive 0; Mismatches 0; Indels
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100.0%; Pred. No. ...
... 0; Mismatches

    .31
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                                                                                                                                                                         137 CCTGCCGCCTTCCACCGTTCATTCTAGAG 165
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GERON CORP (US)
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/mod_base=OTHER
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                                                                                                                                                                                          30 CCTGCCGCCTTCCACCGTTCATCTAGAG 2
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Sequence 2 from Patent WO9845450.
A84592.
A84592.1 GI:6733508
                         1. .30
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Atkinson, E.M. and Kealey, J.T.
PURIFIED TELOMERASE
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          Location/Qualifiers
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Purified telomerase.
BD058133
BD058133.1 GI:22603739
JP 2001509681-A/2.
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Best Local Similarity 100.
Matches 29; Conservative
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unclassified.
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Matches 29; Conserv
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Zea mays
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A84592/c
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopaida; Poales; Poaceae; PACCAD clade; Panicoideae; Andropogoneae; Zea.

I (bases 1 to 30)

Reinrich, S.L., Iii, E.M.A., Lichtsteiner, S.P., Vasserot, A.P., Purizin, S.L., Iii, E.M.A., Lichtsteiner, S.P., Vasserot, A.P., Purizin, A. and Kealey, J.T.

Purizin d telomerase

Patent: JP 2001509681-A/4

PA 24-JUL-2001

PR 04-APR-1997 JP 1998542718

PR 04-APR-1997 JP 1998542718

PI SCOTT L WENENEICH EDWARD A PRUZAN, JAMES T KEALEY PC CIZNIS/54, C12N9/12, C07K16/40, C12Q1/68, C07K14/47 CC Strandedness: Single; CC /mod_base= OTHER

FT modified base

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Weinrich, S.L., Atkinson, E.M. III, Lichtsteiner, S.P., Vasserot, A.P., Pruzan, R.A. and Kealey, J.T.
Purified telomerase
Parent: US 5968506-A 4 19-OCT-1999;
Location/Qualifiers
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                                 PAT 31-AUG-2000
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96.6%; Pred. No. 52;
:ive 0; Mismatches
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Best Local Similarity 96.6%; Pred. No. 52;
Matches 28; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                          /mol_type="unassigned DNA'
                               AR079891 30 bp
Sequence 4 from patent US 5968506.
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/mol_type="genomic DNA"
/db_xref="taxon:4577"
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                                                                                                                                                                                                                                                                           /organism="unknown"
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Purified telomerase.
BD058135
BD058135.1 GI:22603741
JP 2001509681-A/4.
Zea mays
Zea mays
                                                                               AR079891.1 GI:10006644
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Matches 28; Conservative
                                                                                                                               Unknown.
Unclassified.
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RESULT 52
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BD058135/c
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Pred. No. 45;
0; Mismatches 1; Indels
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Atkinson, E.M. and Kealey, J.T.
PURIFIED TELOMERASE
Batent: WO 9845450-A 4 15-OCT-1998;
GERON CORP (US)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /mol_type="unassigned_DNA"
/db_xref="taxon:32644"
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Greider, C.W. and Le, S.
Telomerase-associated proteins
Patent: US 6300131-A 3 09-OCT-2001;
Location/Qualifiers
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/mod_base=OTHER
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/organism="unidentified"
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Sequence 4 from Patent WO9845450.
A84594
/organism="unknown"
/mol_type="genomic DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                             1. .28
/organism="unknown"
/mol_type="genomic DNA"
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                                                ch 6.3%;
1 Similarity 96.7%;
29; Conservative
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A84594/c
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AL Patent: JP 2001507229-A 26 05-JUN-2001;
GERON CORP
PN JP 200150729-A/26
PD 05-JUN-2001
PF 19-DEC-1997 JP 1998529003
PR 20-DEC-1996 US 08/770565 PI
NAM WOO KIM,FRED WU,JAMES T KEALEY,RONALD PRUZAN,SCOTT L PI
WEINRICH
PC C12NN/10, A61K9/08,A61K31/7105,A61K45/00,A61P35/00,PC
C12NN/10,C12Q1/68,C12Q1/68,C12Q1/68,C12N15/00,C12N5/00 CC
Strandedness: Single
CC Topology: Linear;
CC Topology: Linear;
CC Topology: Linear;
FH Key
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CANCER RESEARCH CAMPAIGN TECHNOLOGY LTD
OS Artificial Sequence
RN JP 2002509699-A/4
PD 02-APR-2002
PP 20-JAN-1999 JP 2000529424
PR 29-JAN-1999 GB 9801902.9
PI WILLIAM NICOL KEITH
PC 21-AN-1998 GB 9801902.9
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Promoter region of mouse and human telomerase RNA component genes.
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PC A61P35/00,C12N1/15,C12N1/19,C12N1/21,C12N5/10,C12P21/02
C12Q1/68//C12N9/12,
PC (A61K35/76,A61K31:522),C12N15/00,A61K37/02,C12N5/00 CC
Description of Artificial Sequence: Primer
FH Key Location/Qualifiers
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/organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
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50;
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JP 2002509699-A/4.
synthetic construct
synthetic construct
other sequences; artificial sequences.
1 (bases 1 to 30)
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Best Local Similarity 93.3%; Pred. No. 58;
Matches 28; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                               1. .27
/organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       144 CCTTCCACCGTTCATTCTAGAGCAAAC 170
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BD225801/c
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BD023722.1 GI:22564945
JP 200120229-A/26.
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Kim,N.W., Wu,F., Kealey,J.T., Pruzan,R. and Weinrich,S.L.
Method for detecting and inhibiting RNA component of telomerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Chen, X.Q. and Anker, P.
Cancer diagnosis method
Patent: WO 0190409-A 2 29-NOV-2001;
Chen, Xu Qi (US) ; Stroun, Maurice (CH) ; Anker, Philippe (CH)
Location/Qualifiers
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1 (bases 1 to 27)

1 (hases 1 to 27)

1 (kaley,J.T., Pruzan,R. and Weinrich,S.L. Methods for detecting the RNA component of telomerase Patent: US 5846723-A 26 08-DEC-1998;
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6.0%; Score 27; DB 1; Length 27;
Best Local Similarity 100.0%; Pred. No. 50;
Matches 27; Conservative 0; Mismatches 0; Indels
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/organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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                                                                       27 bp
Sequence 26 from patent US 5846723.
AR063850
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/mol_type="unassigned DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 2 from Patent W00190409.
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synthetic construct
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Matches 2
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AR063850/c
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BD023722/c
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VI OS Unidentified

OS Unidentified

PN JP 2002272489-A/33

PD 24-SEP-2002

PF 06-MAR-2002 UP 2002061125

PF 07-UUL-1995 US 08/272102,27-OCT-1994 US 08/330123 PR

O7-UUL-1995 US 08/272802,07-UNL-1995 US 08/482115 PI BRYANT

VILLEPONTEAU, UTNLI FEW WALTER FUNK, WILLIAM H ANDREWS PC

C12N15/09,C12N9/99,C12Q1/68,G01N33/53,G01N33/566,C12N15/00 CC

Strandedness: Single:

CC Topology: Linear;

CC Mammalian telomerase

FH Key
                                                                                                                                                            Length 28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 28;
                      Andrews, W.H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 (bases 1 to 28)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Wammanlian telomerase
Patent: US 5958680-A 38 28-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               unidentified unclassified.

1 (bases 1 to 28)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H. Mammalian telomerase
Patent: JP 2002272489-A 33 24-SEP-2002;
GERON CORP
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1 (bases 1 to 28)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H Assays for the DNA component of human telomerase Patent: US 577679-A 40 07-JUL-1998;
Location/Qualifiers
                                                                                                                                                            Query Match 5.9%; Score 26.4; DB 1;
Best Local Similarity 96.4%; Pred. No. 57;
Matches 27; Conservative 0; Mismatches 1;
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                                                                                                                                                                                                                                                                                                                                                            DNA
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                                                                                                          /organism="unknown"
/mol_type="unassigned DNA"
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                                                                                                                                                                                                                                  17 GCCTGGGAGGGGTGGTGGCCATTTTTG 44
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JP 2002272489-A/33.
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BD176174
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Best Local Similarity
Matches 27; Conserv
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AR075541
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                                                                                                                                                                                                                                                                                                   Patent: WO 9938964-A 4 05-AUG-1999;
KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH (GB)
Location/Qualifiers
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Unclassified.
1 (bases 1 to 28)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Villeponteau, B., Reng, J., Funk, W. and Andrews, W.H.
Assays for the DNA component of human telomerase
Patent: US 5776679-A 31 07-JUL-1998;
Location/Qualifiers
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/organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="teaxon:32630"
/noTe="primer"
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synthetic construct
other sequences; artificial sequences.
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Best Local Similarity 93.3%; Pred. No. 58;
Matches 28; Conservative 0; Mismatches
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5.9%; Score 26.4; D
Best Local Similarity 96.4%; Pred. No. 57;
Matches 27; Conservative 0; Mismatches
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/organism="unknown"
/mol_type="unassigned DNA"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AR016072 28 bp
Sequence 40 from patent US 5776679.
AR016072
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                                                                                    30 bp
Sequence 4 from Patent WO9938964.
AX019550
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Unclassified.
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PAT 18-MAR-2003

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RESULT 60 AR016072 LOCUS

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PAT 05-DEC-1998
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1 (bases 1 to 26)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H. Assays for the DNA component of human telomerase Patent: US 5776679-A 23 07-JUL-1998;
Location/Qualifiers
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1 (bases 1 to 26)
1 (bases 1 to 26)
1 Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H. Assays for the DNA component of human telomerase Patent: US 5776679-A 29 07-UUL-1998;
Location/Qualifiers
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5.8%; Score 26; DB 1; Length 26;
Best Local Similarity 100.0%; Pred. No. 56;
Matches 26; Conservative 0; Mismatches 0; Indels
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/organism="unknown"
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Sequence 25 from patent US 5858777.
AR028785
AR028785.1 GI:5940758
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Sequence 23 from patent US 5776679.
AR016055
                                                                                                                                                                                                                                                                                                                                    /organism="unknown"
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Sequence 29 from patent US 5776679.
AR016061.1 GI:3972338
                                                                                                                                                                                                                                                                                                                                                                                                                                             145 CTTCCACCGTTCATTCTAGAGCAAAC 170
               145 CTTCCACCGTTCATTCTAGAGCAAAC 170
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                                 26 CTTCCACCGTTCATTCTAGAGCAAAC
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Unclassified.
1 (bases 1 to 26)
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AR028785
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Method for detection of carcinoma of the urinary bladder within a urine sample
Parent: EP 0926245-A 1 30-JUN-1999;
ROCHE DIAGNOSTICS GMBH (DE)
Location/Qualifiers
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Emrich, T.D.
Method for detection of carcinoma of the urinary bladder within a
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ce 1, .28
/organism='Unidentified'.
Location/Qualifiers
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    .26
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    /db_xref="taxon:32644"

    .26
    /organism="unidentified"
    /mol_type="unassigned DNA"
    /db_xref="taxon:32644"

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Patent: EP 0926245-A 2 30-JUN-1999;
ROCHE DIAGNOSTICS GMBH (DE)
Location/Qualifiers
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    .28
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    /mol_type="genomic DNA"
    /db_xref="taxon:32644"

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Sequence 1 from Patent EP0926245.
A94987
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A94987
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PAT 30-AUG-2000
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                                                                                                                                                         Unknow...
Unclassified.
1 (bases 1 to 26)
Kim,N.Woo., Wu,F., Kealey,J.T., Pruzan,R. and Weinrich,S.L.
Kim,N.Woo., Wu,F., Kealey,J.T., Pruzan,R. and Weinrich,S.L.
Methods for detecting the RNA component of telomerase
Patent: US 5846723-A 25 08-DEC-1998;
Location/Qualifiers
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Unclassified.
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Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Mammalian telomerase
Patent: US 5958680-A 24 28-SEP-1999;
Location/Qualifiers
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Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Mammalian telomerase
Patent: US 5958680-A 30 28-SEP-1999;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                     5.8%; Score 26; DB 1; Length 26; llarity 100.0%; Pred. No. 56; Conservative 0; Mismatches 0; Indels
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Sequence 30 from patent US 5958680.
AR075533
AR075533.1 GI:10002281
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/wol_type="unassigned DNA"
                                                                                                                                                                                                                                                                         /organism="unknown"
/mol_type="unassigned DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 24 from patent US 5958680.
AR075527
                                                 AR063849 26 bp
Sequence 25 from patent US 5846723.
AR063849
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Matches 26; Conserv
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AR063849
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AR075527/c
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Unclassified.
I (bases 1 to 26)
Villeponteau, B., Feng, J., Andrews, W.H. and Adams, R.R.
Methods and reagents for regulating telomere length and telomerase activity
Villeponteau, B., Feng, J., Andrews, W.H. and Adams, R.R.
Methods and reagents for regulating telomere length and telomerase
activity
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Villaponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Mammalian telomerase
Patent: US 5837857-A 23 17-NOV-1998;
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Best Local Similarity 100.0%; Prad. No. 56;
Matches 26; Conservative 0; Mismatches
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                                            Patent: US 5858777-A 25 12-JAN-1999;
                                                                                                                                                                                                                                                                                                  26 bp 1
Sequence 26 from patent US 5858777.
AR028786
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/organism="unknown"
/mol_type="unassigned DNA"
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Sequence 23 from patent US 5837857.
AR059216
AR059216.1 GI:5984793
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/organism="unknown"
/mol_type="unassigned DNA"
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/organism="unknown"
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Best Local Similarity 100.
Matches 26; Conservative
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AR028786/c
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Method for detecting bladder cancer in urine samples.
E36507
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Method for detecting bladder cancer in urine samples
Patent: JP 199243995-A 1 14-SEP-1999;
ROCHE DIAGNOSTICS GMBH
OS Artificial Sequence
DN JP 1999243995-A/1
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/organism='Unidentified'
Location/Qualifiers
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synthetic construct
other sequences; artificial sequences.
                                                                                          Query Match 5.8%; Score 26; DB Best Local Similarity 100.0%; Pred. No. 56; Matches 26; Conservative 0; Mismatches
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/organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
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JP 2002272489-A/30.
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E36507
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OS Unidentified

N DP 2002272489-A 25 24-SEP-2002;

OS Unidentified

PD 24-SEP-2002

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(bases 1 to 26)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H. Mammalian telomerase
Patent: US 6258535-A 23 10-JUL-2001;
Location/Qualifiers
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                                                                                                                             Length 26;
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/organism='Unidentified'.
Location/Qualifiers
                                                                                                                          Query Match 5.8%; Score 26; DB 1; Best Local Similarity 100.0%; Pred. No. 56; Matches 26; Conservative 0; Mismatches C
                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA
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                                                              /mol_type="unassigned DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                  26 bp
Sequence 23 from patent US 6258535.
AR161925
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/organism="unknown"
                               /organism="unknown"
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JP 2002272489-A/25.
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BD176166
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AR161925/c
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BD176166/c
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FEATURES

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C12Q1/02,
C12Q1/48, C12Q1/68, G01N33/15, G01N33/48, G01N33/50//C07K14/47, PC
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                                                                                                                                                                                                                                                                                                                   CALVIN B HAREI, WILLIAM H ANDREWS
C12N15/09,A61K31/70,A61K38/55,A61K39/395,A61K39/395,A61K48/00,
                                                                                                                                                                                                                                                                                                                                                                                                                        PC (C12N1/21,C12R1:19),(C12N9/12,C12R1:19),(C12N9/12,C12R1:84),
PC (C12N3/12,C12R1:91),C12N15/00,A61K37/64,C12N5/00 CC
Strandedness: Single;
CC Topology: Linear;
FH Key Location/Qualifiers
FT source Location/Qualifiers
/organism='Unidentified'
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08/844.419,
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08/911.312, PR
08/915.503 PI
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PN JP 199253177-A/254
PD 21-SEP-1999
PF 15-OCT-1996 UP 08/724.643,18-APR-1997 US 08/844.419
PR 01-OCT-1996 US 08/724.643,18-APR-1997 US 08/846.017,06-MAY-1997 US 08/846.017,06-MAY-1997 US 08/813.1312, PR 09-MAY-1997 US 08/812.951,14-AUG-1997 US 08/911.312, PR R SECHI, JOCHIMU RINGNER, TORU NAKAMURA, KAREN B CHAPMAN, PI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  unidentified
unidentified
unidentified
unclassified.
1 (base 1 to 26)
Thomas.R.S., Jochimu.R., Toru,N., Karen,B.C., Greg,B.M.,
Calvin,B.H. and William,H.A.
Human telomerase caralytic subunit promoter
Patent: JP 1999523177-A 254 21-SEP-1999;
JERON CORP, UNIVERSITY TECHNOLOGY CORP
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Human telomerase catalytic subunit promoter.
E37046

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    /organism="unidentified"

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/db_xref="taxon:32644"
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JP 1999253177-A/254.
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AUTHORS
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JOURNAL
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E37046/c
LOCUS
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Method for detecting bladder cancer in urine samples
patent. JP 199924395-A 2 14-SEP-1999;
RCHE DIAGNOSTICS GMBH
OS Artificial Sequence
RN JP 199924395-A/2
PD 14-SEP-1999
PF 22-DEC-1999 JP 1998365689
PF 22-DEC-1997 DE 19757300:2
PI THOMAS ENRIHI
PC C12Q1/68//C12N1/00
CC
FH Key
Location/Qualifiers
FT source
Location/Qualifiers
FT Source
//organism='Artificial Sequence'.
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Method for detecting bladder cancer in urine samples.
E36508
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Location/Qualifiers
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Human telomerase catalytic subunit promoter.
E37045.1 GI:13023008
19799253177-A/253.
                                                                                                          Location/Qualifiers

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synthetic construct
other sequences; artificial sequences.
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100.0%; Pred. No. 56;
tive 0; Mismatches
   14-SEP-1999
22-DEC-1998 JP 1998365689
22-DEC-1997 DE 19757300:2
THOMAS ENRIHI
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                                                                                                                                                                                                                                                                                                                                     54 AACTGAGAAGGGCGTAGGCGCCGTGC 79
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JP 1999243995-A/2.
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Best Local Similarity 100.
Matches 26; Conservative
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Best Local Similarity
Matches 26; Conserve
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ACCESSION VERSION KEYWORDS SOURCE ORGANISM

REFERENCE AUTHORS TITLE

JOURNAL

COMMENT

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DEFINITION

RESULT 77 E36508/c LOCUS

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Gaps

THOMAS GREG B

E37045 LOCUS DEFINITION ACCESSION

VERSION KEYWORDS SOURCE

RESULT 78

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Gaps

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PAT 20-DEC-2002

linear

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1 (bases 1 to 26)
Cech.Y.R., Lingner.J., Nakamura,T., Chapman,K.B., Morin,G.B.,
Harley,C.B. and Andrews,W.H.
Human telomerase catalytic subunit: diagnostic and therapeutic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5.8%; Score 26; DB 1; Length 26; ilarity 100.0%; Pred. No. 56; Conservative 0; Mismatches 0; Indels
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Patent: US 6548298-A 23 15-APR-2003;
Location/Qualifiers
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                                                                                                              Query Match 5.8%; Score 26; DB 1; Length 26; Best Local Similarity 100.0%; Pred. No. 56; Matches 26; Conservative 0; Mismatches 0; Indels
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Location/Qualifiers
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Sequence 312 from patent US 6475789.
AR243519
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Sequence 23 from patent US 6548298.
AR306473
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/organism="unknown"
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/organism="unknown"
    Location/Qualifiers
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Unclassified.
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Matches 26; Conserv
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AR243519/c
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CALVIN B HARBI, WILLIAM H ANDREWS
CI2NIS/09, A61K31/70, A61K38/55, A61K39/395, A61K39/395, A61K48/00,
CI2Q1/02,
CI2Q1/48, CI2Q1/68, G01N33/15, G01N33/48, G01N33/50//C07K14/47, PC
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                                                                                                            C12R1:84),
PC (C12N1/21, C12R5/10, C12N9/12, C12P21/08, (C12N1/19, PC (C12N1/21, C12R1:19), (C12N9/12, C12R1:84),
PC (C12N1/21, C12R1:19), (C12N9/12, C12R1:84),
PC (C12N1/21, C12R1:19), (C12N1/2), (C12N1/21), (
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Cech.T.R., Lingmer.J., Nakamura,T., Chapman,K.B., Morin,G.B.,
Harley,C.B. and Andrews,W.H.
Human telomerase catalytic subunit: diagnostic and therapeutic
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(Millepages 1 to 26)
Villepages 1 to 26)
Villepanteau, B., Feng, J., Funk, W. and Andrews, W.H.
Mammalian telomerase
Patent: US 5583016-A 23 10-DEC-1996;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              5.8%; Score 26; DB 1; Length 26; 100.0%; Pred. No. 56; tive 0; Mismatches 0; Indels
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Pred. No. 56;
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100.0%; Pred. No. co.
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Sequence 311 from patent US 6475789.
AR243518.1 GI:27290729
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Sequence 23 from patent US 5583016.
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/organism="unknown"
/mol_type="unassigned DNA"

    .26
    /organism="unidentified"
    /mol_type="genomic DNA"
    /db_xref="taxon:32644"

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Matches 26; Conservative
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I31770/c
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Kopreski,M.S. and Gocke,C.D.
Method for detection of hTR and hTERT telomerase-associated RNA in plasma or serum
Patent: US 6607988-A 4 19-AUG-2003;
Location/Qualifiers
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Best Local Similarity 100.0%; Pred. No. 56;
Matches 26; Conservative 0; Mismatches
                                            Query Match 5.8%; Score 26; DB Best Local Similarity 100.0%; Pred. No. 56; Matches 26; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                    1 (bases 1 to 26)
Greider, C.W. and Le.S.
Telomerase-associated proteins
Patent: US 6300131-A 4 09-OCT-2001;
Location/Qualifiers
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Sequence 4 from patent US 6300131.
AR370169
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Sequence 5 from patent US 6607898.
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/wol_type="genomic DNA"
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                                                                                                    145 CTTCCACCGTTCATTCTAGAGCAAAC 170
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/wol_type="genomic DNA"
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AR381130/c
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Unclassified.

E 1 (bases I to 26)

S Villeponteau, B., Feng, J., Andrews, W.H. and Adams, R.R.

S Villeponteau, B., Feng, J., Andrews, W.H. and Adams, R.R.

Peptides related to TPC2 and TPC3, two proteins that are coexpressed with teleomerase activity

AL Patent: US 6300110-0.A 26 09-OCT-2001;

Location/Qualifiers
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Unknown.
Unclassified.
I (bases 1 to 26)
Villeponteau, B., Feng, J., Andrews, W.H. and Adams, R.R.
Villeponteau, B., Feng, J., Andrews, W.H. and Adams, R.R.
Peptides related to TPC2 and TPC3, two proteins that are
coexpressed with telomerase activity
Patent: US 6300110-A 25 09-0CT-2001;
Location/Qualifiers
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Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Mammalian telomerase
Patent: US 6548296-A 30 15-APR-2003;
Location/Qualifiers
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56;
           26 bp I Sequence 30 from patent US 6548298.
AR306480
AR306480.1 GI:31696319
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Sequence 25 from patent US 6300110.
AR369722
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AR369723
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/mol_type="genomic DNA"

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    /organism="unknown"
    /mol_type="genomic DNA"

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Unclassified.
Unclassified.
1 (bases 1 to 26)
Cech.T.R., Lingner,J., Nakamura,T., Chapman,K.B., Morin,G.B.,
Harley,C.B. and Andrews,W.H.
Cells immortalized with telomerase reverse transcriptase for use in
                                                                                                                                             PAT 18-DEC-2003
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(bases 1 to 26)

(cch.T.R., Lingmer,J., Nakamura,T., Chapman,K.B., Morin,G.B., Harley,C.B. and Andrews,W.H.
Cells immortalized with telomerase reverse transcriptase for use
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Patent: US 6617110-A 597 09-SEP-2003;
Location/Qualifiers
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Patent: US 6617110-A 598 09-SEP-2003;
Location/Qualifiers
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Sequence 597 from patent US 6617110.
AR393334
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AR393335
Best Local Similarity 100.0%; Pred. No. 56; Matches 0; Mismatches
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Sequence 25 from Patent EP0953042.
AX022186
                                           145 CTTCCACCGTTCATTCTAGAGCAAAC 170
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                                                              Unclassified.

1 (bases 1 to 26)

Kopreski, M. S. and Gocke, C.D.

Method for detection of hTR and hTERT telomerase-associated RNA in plasma or serum

Patent: US 6607898-A 5 19-AUG-2003;

Location/Qualifiers
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Morin,G.B. and Andrews,W.H.
Promoter for telomerase reverse transcriptase
Patent: US 6610839-A 597 26-AUG-2003;
Location/Qualifiers
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Sequence 597 from patent US 6610839.
AR390720
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Sequence 598 from patent US 6610839.
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/organism="unknown"
/mol_type="genomic DNA"
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/organism="unknown"
/mol_type="genomic DNA"
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 AR381130.1 GI:40088891
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Cancer diagnosis method
Patent: WO 0190409-A 1 29-NOV-2001;
Chen, Xu Qi (US) ; Stroun, Maurice (CH) ; Anker, Philippe (CH)
Location/Qualifiers
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Detecting telomerase activity
Patent: WO 0046601-A 9 10-AUG-2000;
LARSEN FRANK (NO); SKAANSENG MARIANNE (NO)
LOCATION/QUALIFIERS
 1. .26
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/db_xref="taxon.32630"
/note="PCR primer"
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Sequence 1 from Patent WO0190409.
AX317988
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                                                                    /note="PCR primer"
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                                                                                                  Andrews,W.H., Villeponteau,B., Adams,R.R. and Feng,J. Methods and reagents for regulating telomere length and telomerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Andrews,W.H., Villeponteau,B., Adams,R.R. and Feng,J.
Methods and reagents for regulating telomere length and telomerase activity
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Detecting telomerase activity
Patent: WO 0046601-A 8 10-AUG-2000;
LARSEN FRANK (NO) ; SKAANSENG MARIANNE (NO)
LOCATION/Qualifiers
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synthetic construct
other sequences, artificial sequences.
                                                                                                                                 activity
Patent: EP 0953042-A 25 03-NOV-1999;
GERON CORP (US)

    .26
    /organism="unidentified"
    /mol_type="unassigned DNA"
    /db_xref="taxon:32644"

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/mol_type="unassigned DNA"
/db_xref="taxon:32644"
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GERON CORP (US)
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Sequence 26 from Patent EP0953042.
AX022187
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Sequence 8 from Patent WO0046601.
AX033376
AX033376.1 GI:10280150
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   AX022186.1 GI:10045854
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N BD011296
BD011296;
BD011296;
JP 201081042-A/253.
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SM unidentified
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S Sechi,T.R., Lingner,J., Nakamura,T., Chapman,K.B., Mori,G.B.,
Harley,C.B. and Andrews,W.H.
                                                                                Cech,T.R., Lingner,J., Nakamura,T., Chapman,K.B., Morin,G.B., Harley,C.B. and Andrews,W.H.
Human telomerase catalytic subunit
Patent: EP 333094-A, 599 06-AUG-2003;
Geron Corporation (US); University Technology Corporation (US)
Location/Qualifiers
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Human telomerase catalytic subunit
Patent: EP 1333094-A 600 60-AUG-2003;
Geron Corporation (US); University Technology Corporation (US)
Location/Qualifiers
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Patent: JP 2001081042-A 253 27-MAR-2001;
GERON CORP,UNIVERSITY TECHNOLOGY CORP
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56;
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/db_xref="taxon:32644"

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Method for detection of htr and htert telomerase-associated rna in plasma or serum
Patent: WO 0218652-A 4 07-MAR-2002;
Oncomedx, Inc. (US)
Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                    Homo sapiens
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.
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Method for detection of htr and htert telomerase-associated rna plasma or serum
Patent: WO 0218652-A 5 07-MAR-2002;
Oncomedx, Inc. (US)
Location/Qualifiers
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/db_xref="taxon:9606"

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/mol_type="unassigned DNA"
/db_xref="taxon:9606"

                                                                                         AX468454 26 bp
Sequence 4 from Patent WO0218652.
AX468454
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Sequence 5 from Patent WO0218652.
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C07K5/10,
                                                                                                                                                                     CALVIN B HARLEY, WILLIAM H ANDREWS
AGIK38/00,AGIK31/7088,AGIK39/00,AGIK48/00,AGIP35/00,AGIP43/00,
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CO7K5/107,CO7K5/117,CO7K7/06,C07K7/08,C07K16/40,C12N9/12, PC
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Sechi,T.R., Lingner,J., Nakamura,T., Chapman,K.B., Mori,G.B.,
Harley,C.B. and Andrews,W.H.
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PC C07K5/10,
PC C07K5/10,C07K5/117,C07K7/06,C07K7/08,C07K16/40,C12N9/12
C12N15/09,
PC C12Q1/02,C12Q1/48,C12Q1/68,G01N33/15,G01N33/50,G01N33/53
G01N33/53,
PC G01N33/56,G01N33/573//C12P21/08,AGIK37/02,C12N15/00 CC Strandedness: Single;
CC Topology: Linear;
                                                           08/844419
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PN JP 2001081042-A/253
PD 27-MAR-2001
PF 27-JUL-2000 JP 2000227474
PR 01-OCT-1996 US 08/724643,18-APR-1997 US 08/84441
25-APR-1997 US 08/846017,06-MAY-1997 US 08/851843 PR 09-MAY-1997 US 08/85050,14-AUG-1997 US 08/913132 PR 14-AUG-1997 US 08/912591,14-AUG-1997 US 08/912591,14-AUG-1997 US 08/912591,14-AUG-1997 US 08/912593 PI R SECHI,JOACHIM LINGNER,TORU NAKAMURA, KAREN B CHAPMAN, PI MORIN,
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.larity 100.0%; Pred. No. 56;
Conservative 0; Mismatches 0; Indels
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Patent: JP 2001081042-A 254 27-MAR-2001;
GERON CORP, UNIVERSITY TECHNOLOGY CORP
                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          26 bp
Human telomerase catalytic subunit.
BD011297
                                                                                                                                                                                                                                                                                                                                                                                                                                                        organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
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JP 201081042-A/254
27-MAR-200
27-JUL-2000 JP 2000227474
01-OCT-1996 US 08/724643
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              45 TCTAACCCTAACTGAGAAGGGCGTAG 70
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/organism='Unidentified'.
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JP 2001081042-A/254.
                                                                                                                                                             PI CALVIN B HARLEY, W)
PC A61K38/00, A61K31/7
PC C07K5/10,
PC C07K5/10,
PC C12Q1/02, C12Q1/48,
PC C12Q1/02, C12Q1/48,
PC G01N33/53,
PC G01N33/56, G01N33/
Strandedness: Single,
CC Topology: Linear;
FH KEY
FT Source 1
FT Aoganism='Unident
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26; Conserva
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26 bp DNA linear PAT 18-SEP-2002 Telomerase assay of bodily fluid for cancer screening and evaluation of disease phase and prognosis.
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Method for detecting and inhibiting RNA component of telomerase.
BD023721
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Kim,N.W., Wu,F., Kealey,J.T., Pruzan,R. and Weinrich,S.L.
Method for detecting and inhibiting RNA component of telomerase
Patent: JP 2001507229-A 25 05-JUN-2001;
GERON CORP
PN JP 2001507229-A/25
PN JP 2001507229-A/25
PN JP 1998529003
PR 20-DEC-1996 US 08/770564,20-DEC-1996 US 08/770565 PI
NAM WOO KIM,FRED WU,JAMES T KEALEY,RONALD PRUZAN,SCOTT L PI
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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PC C12N5/10.

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56;
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JP 2002503480-A/3.
synthetic construct
synthetic construct
other sequences; artificial sequences.
1 (bases 1 to 26)
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Best Local Similarity 100.0%; Pred. No. 56;
Matches 26; Conservative 0; Mismatches
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100.0%; Pred. No. 56;
ative 0; Mismatches
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/db_xref="taxon:32644"

    .26
    /organism="unidentified"

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source 1. .26 /organism='Unidentified'.
                                                                                       Location/Qualifiers
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Page 28

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COTK14/00,A61K38/16,C12Q1/68
Strandedness: Single;
Topology: Linear;
Modulation of mammalian telomerase by peptide nucleic acids FH
Location/Qualifiers
                                                                                                                                                PAT 07-SEP-2000
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Modulation of mammalian telomerase by peptide nucleic acids.
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JP 2001517929-A/39
09-007-2001
09-007-2010
09-APR-1997 JP 1997536487
09-APR-1996 US 08/630019
JERRY W SHAY, WOODRING E WRIGHT, MIECZYSLAW A PIATYSZEK, DAVID
                                                                                                                                                                                                                                                                                                                                Promoter regions of the mouse and human telomerase rna component
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1 (bases 1 to 25)
Shay,J.W., Wright,W.E., Piatyszek,M.A., Corey,D. and Norton,J.C. Modulation of mammalian telomerase by peptide nucleic acids
Patent: JP 2001517929-A 39 09-OCT-2001;
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KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH
Location/Qualifiers
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63;
                                                                                                                                                  linear
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Location/Qualifiers
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/organism="synthetic construct"
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/db_xref="taxon:32630"
/note="primer"
                                                                                                                                                  DNA
                                                                                                                                                                                                                                                                          other sequences; artificial sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5.5%; Score 25; DB 100.0%; Pred. No. 63; ive 0; Mismatches
                                                                                                                                                AX019566 25 bp
Sequence 20 from Patent WO9938964.
AX019566
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/organism="unidentified"
/mol_type="genomic RNA"
/db xref="taxon:32644"
                        175 AATGTCAGCTGCTGGCCCGTTCGCC 199
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                                           25 AATGTCAGCTGCTGGCCCGTTCGCC
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JP 2001517929-A/39.
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JAMES C NORTON
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synthetic construct
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Best Local Similarity 100.
Matches 25; Conservative
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OS Unident
PD 09-0CT-1
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PI JARRY P
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PC CO7K14/
CC TOPOLOG
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ACCESSION
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AX019566/c
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ORGANISM
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DEFINITION
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JOURNAL
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Promoter region of mouse and human telomerase RNA component genes.
BD225817
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                                                                                                                                                                                                                   C12Q1/68,C12N15/09,C12P19/34,C12N15/00
Description of Artificial Sequence: F3b, synthesized, Gibco-BRL
                                                                                                                                          16-FEB-1999 JP 2000531587
16-FEB-1998 US 60/074793
JEFFREX W STROVEL, JUDITH STAMBERG, EDWARD HIGHSMITH, LYNNE V
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PC A61P35/00,C12N1/15,C12N1/19,C12N1/21,C12N5/10,C12P21/02 PC

C1201/68//C12N9/12,

PC (A61K55/76,A61K31:522),C12N15/00,A61K37/02,C12N5/00 CC

Description of Artificial Sequence: Primer
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Strovel,J.W., Stamberg,J., Highsmith,E. and Abruzzo,L.V. Telomerase assay of bodily fluid for cancer screening and evaluation of disease phase and prognosis Patent: JP 2002503480-A 3 05-FBB-2002; UNIVERSITY OF MARYLAND BALTIMORE ON Artificial Sequence PN JP 2002503480-A/3

PD 05-FBB-2002
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Location/Qualifiers
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100.0%; Pred. No. 56;
ive 0; Mismatches 0; Indels
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    .26
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    ^db_xref="taxon:32630"

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    .25
    /organism="synthetic construct"
|mol_type="genomic DNA"
|db_xref="taxon:32630"

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synthetic construct
other sequences; artificial sequences.
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29-JAN-1998 GB 9801902
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Matches 26; Conservative
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BD225817/c
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ORGANISM
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Unclassified.
1 (bases 1 to 28)
1 (lasses 1 to 28)
Willeponteau, B., Feng, J., Funk, W. and Andrews, W.H. Mammalian telomerase
Patent: US 6548298-A 29 15-APR-2003;
Location/Qualifiers
Length 28;
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/organism='Unidentified'.
Location/Qualifiers
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Seguence 29 from patent US 6548298.
AR306479
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Sequence 2 from patent US 5846723.
AR063826.1 GI:5993134
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/db_xref="taxon:32644"
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AR063826/C
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Unclassified.
Unclassified.
1 (bases 1 to 28)
1 villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Assays for the DNA component of human telomerase
Patent: US 5776679-A 28 07-UUL-1998;
Location/Qualifiers
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1 (bases 1 to 28)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H. Mammalian telomerase
Patent: US 5958680-A 29 28-SEP-1999;
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llarity 100.0%; Pred. No. 71;
Conservative 0; Mismatches 0; Indels
                         DB 1; Length 25;
63;
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                       Query Match 5.5%; Score 25; DB Best Local Similarity 100.0%; Pred. No. 63; Matches 25; Conservative 0; Mismatches
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Sequence 28 from patent US 5776679.
AR016060
AR016060.1 GI:3972337
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Mammalian telomerase.
BD176170
BD176170.1 GI:29121876
JP 2002272489-A/29.
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BD176170/c
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AR016060/c
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C12N15/09, A61K31/7105, A61K31/711, A61K35/76, A61K38/00, A61K45/00, PC
                                   Promoter region of mouse and human telomerase RNA component genes Patent: JP 2002509699-A 3 02-APR-2002;
CANCER RESEARCH CAMPAIGN TECHNOLOGY LTD
OS Artificial Sequence
PN JP 2002509699-A/3
PD 02-APR-2002
PP 29-JAN-1999 JP 2000529424
PR 29-JAN-1999 GB 9601902.9
                                                                                                                                                                                                          PC A61F35/00,C12N1/15,C12N1/19,C12N1/21,C12N5/10,C12P21/02 PC A61K35/56 A61K35/00,C12N1/15,C12N1/19,C12N1/21,C12N5/10,C12P21/02 PC CA61K35/76,A61K31:S22),C12N15/00,A61K37/02,C12N5/00 CC Description of Artificial Sequence: Primer FH Key Location/Qualiflers FT source 1.24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1; Length 24; 71;
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Promoter region of mouse and
BD225810
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JP 2002509699-A/13.
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   (bases 1 to 24)
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Method for detecting and inhibiting RNA component of telomerase.
BD023698
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undlassified.
1 (bases 1 to 30)
Kim,N.W., Wu,F., Kealey,J.T., Pruzan,R. and Weinrich,S.L.
Method for detecting and inhibiting RNA component of telomerase
Patent: JP 2001507229-A 2 05-JUN-2001;
GERON CORP
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                                   1 (bases 1 to 30)
Kim, N.Woo., Wu,F., Kealey,J.T., Pruzan,R. and Weinrich,S.L.
Methods for detecting the RNA component of telomerase
Patent: US 5846723-A 2 08-DEC-1998;
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PD 05-JUM-2001
PP 19-DEC-1997 JP 1998529003
PR 20-DEC-1996 US 08/770564,20-DEC-1996 US 08/77056
NAM WOO KIM, FRED WU, JAMES T KEALEY, RONALD PRUZAN, SCOTT L
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PC C12N5/12, C12Q1/68, C12Q1/68, C12N15/00, C12N5/00 CC
Strandedness: Single;
CC Topology: Linear;
CC Topology: Linear;
PH Key
PH Key
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Best Local Similarity 96.2%; Pred. No. 84;
Matches 25; Conservative 0; Mismatches
                                                                                                                                                                 /mol_type="unassigned DNA"
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
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JP 2001507229-A/2.
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                                                                      TITLE
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AUTHORS
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BD225800/c
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PAT 17-JUL-2003
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CANCER RESEARCH CAMPAIGN TECHNOLOGY LTD
CANCER PATE CONTROL OF THE CONTROL OF A LTILICIAL Sequence
NN JP 2002509699-A/13
PD 02-APR-2002
PF 29-JAN-1999 JP 2000529424
PR 29-JAN-1998 GB 9801902.9
PI WILLIAM NICOL KEITH
PC C12N15/09, A61K31/7105, A61K31/711, A61K35/76, A61K38/00, A61K45/00, PC
bp DNA linear PAT 17-JUL-200
human telomerase RNA component genes.
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PC AGIR35/00,CI2N1/15,CI2N1/19,CI2N1/21,CI2N5/10,CI2P21/02

C12Q1/68//CI2N9/12,

PC (AGIX35/76,AGIK31:522),CI2N15/00,AGIK37/02,CI2N5/00 CC
Description of Artificial Sequence: Primer
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    .24
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Indels

Length 24;

PAT 20-DEC-2002

linear

DNA

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Promoter regions of the mouse and human telomerase rna component
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Silverman, R.H., Kondo, S., Cowell, J.K., Li, G. and Torrence, P.F.
RNase L activators and antisense oligonucleotides effective to
treat telomerase-expressing malignancies
Patent: US 6468983 + 5 22-00T-2002;
Location/Qualifiers
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KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH (GB)
Location/Qualifiers
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                                    DB 1;
71;
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                                    5.3%; Score 24; DB ilarity 100.0%; Pred. No. 71; Conservative 0; Mismatches
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Sequence 5 from patent US 6468983.
AR241178
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           423 CGTGCACCCAGGACTCGGCTCACA 446
                                                                                                                             1 TTTGTCTAACCCTAACTGAGAAGG 24
                                                                                                           41 TTTGTCTAACCCTAACTGAGAAGG 64
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AR241178/c
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AX019549/c
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Promoter region of mouse and human telomerase RNA component genes. BD25816.1 GI:3303586
BD225816.1 GI:3303586
JP 2002509599-A/19.
Synthetic construct
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CANCER RESEARCH CAMPAIGN TECHNOLOGY LTD
OS Artificial Sequence
N JP 2002509699-A/19
PD 02-APR-2002
PP 29-JAN-1999 JP 2000529424
PR 29-JAN-1999 GB 9801902.9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C12N15/09, A61K31/7105, A61K31/711, A61K35/76, A61K38/00, A61K45/00, PC
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Silverman, R.H., Kondo, S., Cowell, J.K., Li, G. and Torrence, P.F.
RNase L activators and antisense oligonucleotides effective to
treat telomerase-expressing malignancies
Patent: US 6468983-A 4 22-OCT-2002;
Location/Qualifiers
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PC A61P35/00, C12N1/15, C12N1/19, C12N1/21, C12N5/10, C12P21/02
PC (A61R35/76, A61K31:522), C12N15/00, A61K37/02, C12N5/00 CC
Description of Artificial Sequence: Primer
FH Key 1.24
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/organism='Artificial Sequence'.
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Location/Qualifiers
5.3%; Score 24; DB 1; Length 24;
llarity 100.0%; Pred. No. 71;
Conservative 0; Mismatches 0; Indels
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Sequence 4 from patent US 6468983.
AR241177
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29-JAN-1998 GB 9801902.9
WILLIAM NICOL KEITH
                                                                        46 CTAACCCTAACTGAGAAGGGCGTA 69
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24; Conservative
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 Query Match
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AR241177
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TITLE
JOURNAL
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Gaps

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PAT 07-SEP-2000

linear

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Modulation of mammalian telomerase by peptide nucleic acids.

BD071058 1 G1:22616661

JP 2001517929-A/24.
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1 (bases 1 to 24)

S Shay,J.W., Wright,W.E., Piatyszek,M.A., Corey,D. and Norton,J.C.
Modulation of mammalian telomerase by peptide nucleic acids
GERON CORP

OB Unidentified

N JP 2001517929-A 24 09-OCT-2001;
OS Unidentified

PN JP 2001517929-A/24

PD 09-OCT-2001

PR 09-APR-1997 JP 1997536487

PR 09-APR-1996 US 08/630019

PI JERRY W SHAY,WOODRING E WRIGHT,MIECZYSLAW A PIATYSZEK,DAVID PI JAMES C NORTON

PI JAMES C NORTON

CC COTMIL4/00,AGIX8/16,C12Q1/68

CC Topology: Linear;

CC Topology: Linear;
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linkages are replaced by N-(2-aminoethyl)glycine units linked
                     Au,J.L. and Wientjes,G. Compositions active in telomere damage comprising a taxane and telomerase inhibitor Patent: WO 0074667-A 5 14-DEC-2000;
Au, Jessie L.S. (US); Wientjes, Guillaume (US)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                phosphate

CC linkages are replaced by N-(2-aminoethyl)glycin

CC c nucleotide bases via glycine amino N through a

methylenecarbonyl linker'

FH Key Location/Qualifiers

FT source 1.24

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    .24
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                                                                                                                                                                                                                                                                                                      162 AGAGCAAACAAAAATGTCAGCTG 185
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                                                                                                                                                                     Keith, W.N. Promoter regions of the mouse and human telomerase rna component
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KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH (GB)
Location/Qualifiers
                                                                                                                                                                                                                      Patent: WO 9938964-A 13 05-AUG-1999;
KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH
Location/Qualifiers
                    linear
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synthetic construct
other sequences; artificial sequences.
              Sequence 13 from Patent WO9938964. AX019559
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Sequence 5 from Patent WO0074667,
AX058270
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synthetic construct
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Gaps

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BD131326 25 bp DNA linear PAT 18-SEP-2002 Telomerase assay of bodily fluid for cancer screening and evaluation of disease phase and prognosis.
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Description of Artificial Sequence: R3c, synthesized, G1bco-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16-FEB-1999 JP 2000531587
16-FEB-1998 US 60/074793
JEFFREY W STROVEL, JUDITH STAMBERG, EDWARD HIGHSMITH, LYNNE V
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JP 2002503480-A/4.

Synthetic construct
SM synthetic construct
cother sequences; artificial sequences.

Cother sequences; artificial sequences.

S Strovel,J.W., Stamberg,J., Highsmith,E. and Abruzzo,L.V.
Telomerase assay of bodily fluid for cancer screening and evaluation of disease phase and prognosis
LNIVERSITY OF MARYLAND BALTIMORE
OS Artificial Sequence
PN JP 2002503480-A/4
PD 05-FEB-2002
PP 16-FEB-1999 US 60/074793
PP 16-FEB-1999 US 60/074793
PP JEFRER W STROVEL,JUDITH STAMBERG, EDWARD HIGHSMITH, LYNNE
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Modulation of mammalian telomerase by peptide nucleic acids.
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/organism='Artificial Sequence'
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/organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
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/organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
                                                                                        DB 1;
71;
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                                                                                        5.3%; Score 24; DB 100.0%; Pred. No. 71; iive 0; Mismatches
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                                                                                                                                                                  423 CGTGCACCCAGGACTCGGCTCACA 446
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JP 2001517929-A/25.
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1 (bases 1 to 23)
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                 BD084640 21-AUG-2002 RNase L activators and antisense oligonucleotides effective to treat telomerase-expressing malignancies.
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Synthetic construct
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other sequences; artificial sequences.

E 1 (bases 1 to 24)
S Silverman, R. H., Kondo, S., Cowell, J. K., Li, G. and Torrence, P. F.
RNase Lactivators and antisense oligonucleotides effective to
treat telomerase-expressing malignancies
L Patent: JP 2001524100-A 5 27-NOV-2001;
THE CLEVELAND CLINIC FOUNDATION, NATIONAL INSTITUTES OF HEALTH
OS Artificial Sequence
PN JP 2001524100-A/5
                                                                                                                                                                                                                                                                                              OS Artificial Sequence
PN JP 2001524100-A/4
PD 27-NOV-2001
PF 13-APR-1999 US 60/044507,03-FEB-1998 US 09/018125 PI
ROBERT H SILVERMAN, SEIJI KONDO, JOHN K COWELL, GUIYING LI, PAUL F
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PN JP 2001524100-A/5
PD 27-NOV-2001
PF 13-APR-1998 JP 1998546125
PR 21-APR-1997 US 60/044507,03-FEB-1998 US 09/018125 PI
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other sequences; artificial sequences.

1 (bases 1 to 24)
Silverman, R.H., Kondo, S., Cowell, J.K., Li, G. and Torrence, P.F.
RNase L activators and antiense oligonucleotides effective to
treat telomerase-expressing malignancies
Patent: JP 201524100-A 4 27-NOV-2001;
THE CLEVELAND CLINIC FOUNDATION, NATIONAL INSTITUTES OF HEALTH
PN JP 2001524100-A/4
PN JP 2001524100-A/4
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100.0%; Pred. No. 71;
cive 0; Mismatches 0; Indels
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C07H21/00,C07H21/02,C12Q1/68,A61K48/00
Description of Artificial Sequence: primer
Key
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Description of Artificial Sequence: primer
Key

    .24
    Organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"

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JP 2001524100-A/4.
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PAT 07-SEP-2000
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KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH
Location/Qualifiers
          Length 25;
                                           Indels
                                                                                                                                                                                            linear
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4.9%; Score 22; DB 1; Length 22;
Best Local Similarity 100.0%; Pred. No. 90;
Matches 22; Conservative 0; Mismatches 0; Indels
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/organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="taxon:32630"
/note="Oligonucleotide"
        5.1%; Score 23; DB 1;
100.0%; Pred. No. 87;
tive 0; Mismatches (
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Villeponteau, B., Feng, J., Funk, W. and Assays for the DNA component of human Patent: US 5776679-A 32 07-UUL-1998;
Location/Qualifiers
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AR055220/c
LOCUS
DEFINITION Sequence 27 from patent US 5837857.
                                                                                                                                                                                         AX019590 25 bp
Sequence 44 from Patent WO9938964.
AX019590
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/organism="unknown"
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                                                                                          GGGTTGCGGAGGGTGGGCCTGGG
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synthetic construct
     Query Match
Best Local Similarity 100.0
Matches 23, Conservative
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AR016064/c
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Promoter region of mouse and human telomerase RNA component genes.
BD225841
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                                                                                                                                                                                  Topology: Linear; /desc = 'peptide nucleic acid (PNA), where (deoxy(ribose- CC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Promoter region of mouse and human telomerase RNA component genes Patent: JP 2002509699-A 44 02-APR-2002; CANCER RESEARCH CAMPAIGN TECHNOLOGY LTD SQUENCE ALIficial Sequence PN JP 2002509699-A/44
                                                                09-APR-1997 JP 1997536487
09-APR-1996 US 08/630019
JERRY W SHAY, WOODRING E WRIGHT, MIECZYSLAW A PIATYSZEK, DAVID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A61P35/00,C12N1/15,C12N1/19,C12N1/21,C12N5/10,C12P21/02 PC
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PC A61P35/00,C12N1/15,C12N1/19,C12N1/21,C12N5/10,C12P21,
C12Q1/68//C12N9/12,
PC (A61K35/76,A61K31:522),C12N15/00,A61K37/02,C12N5/00
Description of Artificial Sequence:Oligonucleotide FH K
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FH Key Location/Qualifiers
FT source 1. .23
FT /organism='Unidentified'.
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Location/Qualifiers
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80;
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Location/Qualifiers
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/organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
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JP 2002509699-A/44.
synthetic construct
synthetic construct
other sequences; artificial sequences.
1 (bases 1 to 25)
Keith, W.N.
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JAMES C NORTON
CO7K14/00,A61K38/16,C12Q1/68
Strandedness: Single;
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/organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
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29-JAN-1998 GB 9801902
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                                 JP 2001517929-A/25
09-0CT-2001
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                 Unidentified
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Matches 23; Conservative
GERON CORP
                   Query Match
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Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Mammalian telomerase
Patent: US 5958680-A 42 28-SEP-1999;
Location/Qualifiers
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Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Mammalian telomerase
Patent: US 5958680-A 43 28-SEP-1999;
Location/Qualifiers
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Sequence 42 from patent US 5958680.
AR075545.
AR075545.1 GI:10002291
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4.9%; Score 22; DB
Best Local Similarity 100.0%; Pred. No. 90;
Matches 22; Conservative 0; Mismatches
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Sequence 43 from patent US 5958680.
AR075546.
AR075546.1 GI:10002292
    Mismatches
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/mol_type="unassigned DNA"
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/organism="unknown"
/mol_type="unassigned DNA"
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                           183 CTGCTGGCCCGTTCGCCCTCC 204
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                                            CIGCIGGCCCGIICGCCCCICC 1
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Mammalian telomerase.
BD176150
BD176150.1 GI:29121854
JP 2002272489-A/9.
unidentified
unidentified
unclassified.
    Conservative
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Unclassified.
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LOCUS
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AR075545/c
LOCUS
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AR075546/c
LOCUS
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Unclassified.
Unclassified.
(I (bases 1 to 22)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Mammalian telomerase
Patent: US 5837857-A 28 17-NOV-1998;
Locarion/Qualifiers
                                                    Unknown.
Unclassified.
1 (bases 1 to 22)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H. Mammalian telomerase
Patent: US 5837857-A 27 17-NOV-1998;
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Unknown.
Unclassified.
1 (bases 1 to 22)
1 (bases 1 to 22)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Mammalian telomerase
Patent: US 5958680-A 37 28-5EP-1999;
Location/Qualifiers
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Best Local Similarity 100.0%; Pred. No. 90;
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4.9%; Score 22; DB 1;
Best Local Similarity 100.0%; Pred. No. 90;
Matches 22; Conservative 0; Mismatches
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/organism="unknown"
/mol_type="unassigned DNA"
                                                                                                                                                                                                                                                                                                                                                Sequence 28 from patent US 5837857.
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AR075540
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/organism="unknown"
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/organism="unknown"
/mol_type="unassigned DNA"
                                                                                                                                    Location/Qualifiers
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                GI:5984797
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Best Local Similarity
Matches 22; Conserva
  AR059220
AR059220.1
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AR075540/c
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OS Unidentified

PN JP 2002272489-A/32

PD 24-SEP-2002

PP 24-SEP-2002

PP 06-MAR-2002

PR 07-JUL-1994 US 08/272102,27-OCT-1994 US 08/330123 PR

O7-JUL-1995 US 08/472802,07-JUL-1995 US 08/482115 PI BRYANT

VILLEPONTEAL, JUNIA FENC, WALTER FUNK, WILLIAM H ANDREWS PC

C12N15/09, C12N9/99, C12Q1/68, G01N33/53, G01N33/566, C12N15/00 CC

Strandedness: Single,

CC Topology: Linear;

CC Mammalian telomerase

FH Key Location/Qualifiers

FT source /organism='Unidentified'.
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1 (bases 1 to 22)
Villaponteau, B., Feng, J., Funk, W. and Andrews, W.H. Mammalian telomerase
Patent: JP 2002272489-A 32 24-SEP-2002;
GERON CORP
                                                                                                                                      linear
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Unclassified.
Unclassified.
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Mammalian telomerase
Patent: US 5583016-A 5 10-DEC-1996;
Locarion/Qualifiers
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Pred. No. 90;
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100.0%; Pred. No. 90;
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4.9%; Score 22; DB
Best Local Similarity 100.0%; Pred. No. 90;
Matches 22; Conservative 0; Mismatches
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/organism="unknown"
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Sequence 5 from patent US 5583016.
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/organism="unidentified"
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                             AACTGAGAAGGGCGTAGGCGCC
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JP 2002272489-A/32.
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Best Local Similarity 100.
Matches 22; Conservative
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OS Unidentified

N JP 202272489-A 10 24-SEP-2002;

OS Unidentified

N JP 202272489-A/10

DD 24-SEP-2002

PF 06-MAR-2002 JP 2002061125

PF 06-MAR-2002 JP 2002061125

PF 07-JUL-1994 US 08/330123 PR 07-JUN-1995 US 08/432102, Z7-OCT-1994 US 08/43213 PI BRYANT O7-JUN-1995 US 08/472802, O7-JUN-1995 US 08/472802, O7-JUN-1995 US 08/482115 PI BRYANT C12N15/09.C12N9/99, C12Q1/68,G01N33/53,G01N33/56,C12N15/00 CC Strandedness: Single;

CC Topology: Linear;

CC Mammalian telomerase

FH Key

Location./

FT source
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32115 PI BRYANT
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OE Unidentified

PN JP 200272489-A/9

PP 24-SEP-2002

PP 24-SEP-2002

PF 06-WAR-2002

OB/272102,27-OCT-1994 US 08/330123 PR

O7-UUL-1995 US 08/472802,07-UUL-1995 US 08/482115 PI BRYAN

OT-UUN-1995 US 08/472102,27-OCT-1994 US 08/482115 PI BRYAN

VILLEPONTEAU, UUNLI FENG, WALLTER FUNK, WILLIAM H ANDEWS PC

C12N15/09,C12N9/99,C12Q1/68,G01N33/53,G01N33/566,C12N15/00 CC

CTOPOLOGY: Linear;

CC Mammalian telomerase

FH Key

Location/Qualifiers

FT source

/organism='Unidentified'.
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 1 (bases 1 to 22)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H. Mammalian telomerase
Patent: JP 2002272489-A 9 24-SEP-2002;
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/organism='Unidentified'
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    .22
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JP 2002272489-A/10.
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Mammalian telomerase.
BD176151
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BD176151/c
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and Pruzan,R.A.
Methods for purifying telomerase
Patent: US 654513-A 2 08-APR-2003;
Location/Qualifiers
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Unclassified.
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AR306489/c
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Weinrich,S.L., Atkinson,E.M. III, Lichtsteiner,S.P., Vasserot,A.P. and Pruzan,R.A.
Purified telomerase
Patent: US 6517834-A 2 11-FEB-2003;
Location/Qualifiers
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Unclassified.
1 (bases 1 to 22)
Weinrich, S.L., Atkinson, E.M. III, Lichtsteiner, S.P., Vasserot, A.P.
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Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Mammalian telomerase
Patent: US 5583016-A 6 10-DEC-1996;
Location/Qualifiers
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1 Similarity 100.0%; Pred. No. 90;
22; Conservative 0; Mismatches 0; Indels
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100.0%; Pred. No. 90;
iive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                            /organism="unknown"
/mol_type="unassigned DNA"
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                                                                        131753 22 bp
Sequence 6 from patent US 5583016.
131753.1 GI:1822544
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22 CTAACCCTAACTGAGAAGGGCG 1
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Best Local Similarity 100.
Matches 22; Conservative
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AR305068/c
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AR279617/c
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1 (Dases 1 to 22)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H. Mammalian telomerase
Patent: US 6548298-A 41 15-APR-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (bases 1 to 22)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H. Wammahian telomerase
Patent: US 6548299-A 42 15-APR-2003;
                                                                                                                                                                                                                                        linear
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90;
                                                   DB 1;
                                                 Query Match 4.9%; Score 22; DB Best Local Similarity 100.0%; Pred. No. 90; Matches 22; Conservative 0; Mismatches
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4.9%; Score 22; DB
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Matches 22; Conservative 0; Mismatches
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4.9%; Score 22; DB
Best Local Similarity 100.0%; Pred. No. 90;
Matches 22; Conservative 0; Mismatches
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Sequence 42 from patent US 6548298.
AR306490
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OS Unidentified
PN JP 200227489-A/27
PD 24-SEP-2002
PF 06-WAR-2002 JP 2002061125
PF 06-WAR-2002 JP 2002061125
PF 07-JUL-1994 US 08/272102,27-OCT-1994 US 08/330123 PR 07-JUL-1995 US 08/472802,07-JUN-1995 US 08/482115 PI BRYANT VILLEPONTEAL, UNMI. FENG, WALTER FUNK, WILLIAM HANDREWS PC C12N15/09, C12N9/99, C12Q1/68, G01N33/53, G01N33/566, C12N15/00 CC CT TOPOLOgy: Linear; CC Mammalian telomerase CC Mammalian telomerase FT source Location/Qualifiers
FT source /organism='Unidentified'.
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Unclassfiled.
1 (bases 1 to 21)
Villeponteau B., Feng, J., Funk, W. and Andrews, W.H.
Mammalian telomerata.
Patent: US 625635-A 25 10-JUL-2001;
Location/Qualifiers
                              4.7%; Score 21; DB 1; Length 21; 100.0%; Pred. No. 1e+02;
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llarity 100.0%; Pred. No. 1e+02;
Conservative 0; Mismatches 0; Indels
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Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H. Mammalian telomerase
Patent: JP 2002272489-A 27 24-SEP-2002;
GERON CORP
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/organism='Unidentified'
Location/Qualifiers
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                                         larity 100.0%; Pred. No. 1e-
Conservative 0; Mismatches
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Sequence 25 from patent US 6228535.
AR161927
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    .21
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Mammalian telomerase.
BD176168.1
JP 2002272489-A/27.
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Matches 21; Conserv
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TITLE
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Unclassified.
I (bases It o 21)
I (lasses It o 21)
Assays for the DNA component of human telomerase
Patent: US 5776679-A 25 07-UUL-1998;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unknown.
Unklown.
Unclassified.
Unclassified.
1 (bases 1 to 21)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Mammalian telomerase
Patent: US 5827857-A 25 17-NOV-1998;
Location/Qualifiers
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1 (bases 1 to 21)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Patent: US 5958680-A 26 28-SEP-1999;
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Sequence 25 from patent US 5837857.
AR059218
                           AR016057 21 bp Sequence 25 from patent US 5776679.
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Sequence 26 from patent US 5958680.
AR075529
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Best Local Similarity 100.0
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PAT 31-AUG-2000

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PAT 27-AUG-2002

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Unclassified.

1 (bases 1 to 22)
Weinrich, S.L., Atkinson, E.M. III, Lichtsteiner, S.P., Vasserot, A.P.,
Weinrich, S.A. and Kealey, J.T.
Purified telomerase
Patent: US 5968506-A 3 19-OCT-1999;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; PACCAD calde; Panitooleae; Andropogoneae; Zea.

1 (bases 1 to 22)
Weinrich,S.L., Iii,E.M.A., Lichtsteiner,S.P., Vasserot,A.P., Pruzan,R.A. and Kealey,J.T.
Pruzan,R.A. and Kealey,J.T.
Parent: JP 2001509681-A 3 24-JUL-2001;
GERON CORP
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1 (bases 1 to 22)
Atkinson, E.M. and Kealey, J.T.
PUNFIED TELOMERASE
PATENT TELOMERASE
PATENT WO 9845450-A 3 15-OCT-1998;
GERON CORP (US)
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Sequence 3 from patent US 5968506.
AR079890
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JP 2001509681-A/3.
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Purified telomerase.
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BD058134/c
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AR079890/c
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Unclassified.
Unclassified.
(1 (bases 1 to 21)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H. Mammalian telomerase
Patent: US 6548298-A 25 15-APR-2003;
Location/Qualifiers
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Villeponteau B., Feng,J., Funk,W. and Andrews,W.H.
Mammalian telomerase
Patent: US 5583016-A 25_10-DEC-1996;
                    Length 21;
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                  Query Match 4.7%; Score 21; DB 1;
Best Local Similarity 100.0%; Pred. No. 1e+02;
Matches 21; Conservative 0; Mismatches 0
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Best Local Similarity 100.0%; Pred. No. 1e+02;
Matches 21; Conservative 0; Mismatches 0
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131772
131772.1 GI:1822563
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Sequence 25 from patent US 6548298.
AR306475.1 GI:31696314
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Sequence 3 from Patent WO9845450.
A84593.1 GI:6733509

    .21
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    /mol_type="genomic DNA"

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                                                                                                                                                                                                                                                                                                                                Unknown.
Unclassfiled.
1 (bases 1 to 20)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Mammalian telomerase
Patent: US 5837857-A 7 17-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unclassified.

1 (bases 1 to 20)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H. Mammalian telomerase
Patent: US 5837857-A 26 17-NOV-1998;
Location/Qualifiers
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4.4%; Score 20; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 1.1e+02;
Matches 20; Conservative 0; Mismatches 0; Indels
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                                                          Query Match
4.4%; Score 20; DB 1; Lv
Best Local Similarity 100.0%; Pred. No. 1.1e+02;
Matches 20; Conservative 0; Mismatches 0;
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Sequence 29 from patent US 5837857.
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Sequence 26 from patent US 5837857.
AR059219.1 GI:5984796
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Sequence 7 from patent US 5837857.
AR059200
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                                                                                                                       2 GGTTGCGGAGGGTGGGCCTG 21
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Best Local Similarity
Matches 20; Conserv
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AR059219/c
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AR059200/c
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PN JP 2001509681-A/3
PD 24-JUL-2001
PF 04-ARPL-1997 JP 1998542718
PI SCOTT L WEINRICH EDWARD M ATKINSON III.SERGE P LICHTSTEINER,
PI ALAIN P VASSEROT, RONALD A PRUZAN, JAMES T KEALEY PC
C12N15/54, C12N9/12, C07K16/40, C12Q1/68, C07K14/47 CC Strandedness:
Single;
CC Topology: Linear;
CC /mod_base= OTHER
CC /mod_base= OTHER
CC /note= 'N = biotinylated C'
CC /note= 'Oligonucleotide P3'
FH Key Location/Qualifiers
FT modified base 1:
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1 (bass 1 to 20)
Willeponteau, B., Feng, J., Funk, W. and Andrews, W.H. Assays for the DNA component of human telomerase Patent: US 5776679-A 7 07-JUL-1998;
Location/Qualifiers
                                                                                                                                                                                                                                                                                              Query Match 4.7%; Score 21; DB 1; Length 22; Best Local Similarity 100.0%; Pred. No. 1e+02; Matches 21; Conservative 0; Mismatches 0; Indels
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100.0%; Pred. No. 1.1e+02;
Live 0; Mismatches 0;
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Atkinson, E.M. and Kealey, J.T.
PURIFIED TELOMERASE
PATENT: WO 9845450-A 14 15-OCT-1998;
GERON CORP (US)

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Sequence 7 from patent US 5776679.
AR016039.1 GI:3972316
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Sequence 14 from Patent WO9845450.
A84604
                                                                                                                                                                                                                                                    /mol_type="genomic DNA"
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AR016039/c
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Unknown.
Unclassified.
1 (bases 1 to 20)
Xim,N.Woo., Wu,F., Kealey,J.T., Pruzan,R. and Weinrich,S.L.
Kim,N.Woo. deceting the RNA component of telomerase
Patent: US 5846723-A 13 08-DEC-1998;
Location/Qualifiers
                                                                                                                                                                                                                                         (bases 1 to 20)

Kim,N.Woo., Wu,F., Kealey,J.T., Pruzan,R. and Weinrich,S.L.

Methods for detecting the RNA component of telomerase

Patent: US 5846723-A 7 08-DEC-1998;

Location/Qualifiers
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100.0%; Pred. No. 1.1e+02;
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Best Local Similarity 100.0%; Pred. No. 1.1e+02;
Matches 20; Conservative 0; Mismatches 0;
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                                                                                                                       20 bp
Sequence 7 from patent US 5846723.
AR063831.1 GI:5993139
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Sequence 8 from patent US 5958680.
AR075511
                            300 GAAGAGTTGGGCTCTGTCAG 319
                                                                                                                                                                                                                                                                                                                                                                                                                                  290 CTGCCACCGCGAAGAGTTGG 309
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                                                       GAAGAGTTGGGCTCTGTCAG 1
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AR063831/c
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AR075511/c
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AR063837/c
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AUTHORS
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Unclassified.
Unclassified.
Kim,N. Woo., Wu.F., Kealey,J.T., Pruzan,R. and Weinrich,S.L.
Kim,N. Woo., Wu.F., Kealey,J.T., Pruzan,R. and Weinrich,S.L.
Methods for detecting the RNA component of telomerase
Patent: US 5846723-A 3 08-DEC-1998;
Location/Qualifiers
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Kim,N.Woo., Wu,F., Kealey,J.T., Pruzan,R. and Weinrich,S.L.

Methods for detecting the RNA component of telomerase

Patent: US 5846723-A 6 08-DEC-1998;

Location/Qualifiers
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                                                                                1 (bases 1 to 20)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Mammalian telomerase
Patent: US 5837857-A 29 17-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4.4%; Score 20; DB 1; Length 20; ilarity 100.0%; Pred. No. 1.1e+02; Conservative 0; Mismatches 0; Indels
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Sequence 3 from patent US 5846723.
AR063827.1 GI:5993135
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20 bp
AR063830 GI:5993138
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              GI:5984799
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AR059222
AR059222.1
                                         Unknown.
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AR063827/c
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AR063830/c
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PAT 17-0CT-2001

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PAT 18-MAR-2003

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OS Unidentified

DE 202272489-A/8

PN JP 2002272489-A/8

PD 24-SEP-2002

PF 06-MAR-2002 JP 2002061125

PR 07-JUL-1994 US 08/330123 PR

OF-JUL-1995 US 08/472802,07-JUN-1995 US 08/482115 PI BRYANT

VILLEPONTEAU, JUNIA FENG, WALTER FUNK, WILLIAM H ANDREWS PC

C12N15/09, C12N9/99, C12Q1/68, G01N33/53, G01N33/56, C12N15/00 CC

Strandedness: Single;

CC Mammalian telomerase

FH Key L. 20

FT source Location/Qualifiers

FT F Source Lorganism='Unidentified'.
                                                                                                                                                  Unknown.
Unclassified.
Unclassified.
1 (bases 1 to 20)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Mammalian telomerase
Patent: US 625835-A 7 10-JUL-2001;
Locarion/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 (bases 1 to 20)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H. Mammalian telomerase
Patent: JP 2002272489-A 8 24-SEP-2002;
GERON CORP
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                                                             linear
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                                                            DNA
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                                                     20 bp
Sequence 7 from patent US 6258535.
AR161909
AR161909.1 GI:16228923
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JP 2002272489-A/8.
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Best Local Similarity 100..
Fest 20; Conservative
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AR161909/c
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BD176149/c
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AUTHORS
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Purified telomerase
Parent: US 5968506-A 7 19-OCT-1999;
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1 (bases 1 to 20)
Villeponteau, B., Ferg, J., Funk, W. and Andrews, W.H.
Mammalan telomerase
Patent: US 558680-A 41 28-SEP-1999;
Location/Qualifiers
1 (bases 1 to 20)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Mammalian telomeraepe
Patent: US 5958680-A 8 28-SEP-1999;
Location/Qualifiers
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                                                                                                            /mol_type="unassigned DNA"
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Sequence 41 from patent US 5958680.
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Sequence 7 from patent US 5968506.
AR079894

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Unclassified.
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PAT 17-JUL-2003
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Promoter region of mouse and human telomerase RNA component genes.
BD225842
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CANCER RESEARCH CAMPAIGN TECHNOLOGY LTD
OS Artificial Sequence
PN JP 2002509699-A/45
PP 02-APR-2002
PF 29-JAN-1999 JP 2000529424
PR 29-JAN-1999 GB 9801902.9
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PC A61P35/00,C12N1/15,C12N1/19,C12N1/21,C12N5/10,C12P21/02 PC
,C12Q1/68//C12N9/12,
PC (A61K35/76,A61K31:522),C12N15/00,A61K37/02,C12N5/00 CC
Description of Artificial Sequence:Oligonucleotide FH Key
Location/Qualifiers
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            ce 1. 20 /organism='Artificial Sequence'
Location/Qualifiers
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Location/Qualifiers
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100.0%; Pred. No. 1.1e+02;
ative 0; Mismatches 0; Indels
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    ^mol_type="genomic DNA"
    ^db_xref="taxon:32630"

Location/Qualifiers

    .20
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|mol_type="genomic DNA"
|db_xref="taxon:32630"

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synthetic construct
other sequences; artificial sequences.
1 (bases 1 to 20)
Keith, W.N.
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Sequence 4 from patent US 5583016.
131751
131751.1 GI:1822542
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JP 2002509699-A/45.
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Matches 20; Conserv
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131751/c
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PN JP 200227469-A/11
PD 24-SEP-2002 JP 2002061125
PF 06-MAR-2002 JP 2002061125
PF 07-JUL-1994 US 08/330123 PR 07-JUL-1995 US 08/330123 PR O7-JUL-1995 US 08/482115 PI BRYANT VILLEPONTEAU, JUNID FENG, WALTER FUNK, WILLIAM H ANDREWS PC C12NI5/09, C12N9/99, C12Q1/68, G01N33/53, G01N33/566, C12N15/00 CC Strandedness: Single;
CC Topology: Linear;
CC Mammalian telomerase
FH Key Location/Qualifiers
FT source //organism='Unidentified'.
                                               PAT 18-MAR-2003
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human telomerase RNA component genes.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Promoter region of mouse and human telomerase RNA component genes Patent: JP 2002509699-A 15 02-APR-2002;
CANCER RESEARCH CAMPAIGN TECHNOLOGY LTD
OS ARTIficial Sequence
PN JP 2002509699-A/15
PP 29-JAN-1999 JP 200529424
PR 29-JAN-1996 GB 9801902.9
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PC (A61K35/76,A61K1.522),C12N15/00,A61K37/02,C12N5/00 CC Description of Artificial Sequence: Primer
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Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H. Mammalian telomerase
Patent: UP 2002272489-A 11 24-SEP-2002;
GERON CORP
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Location/Qualifiers
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ilarity 100.0%; Pred. No. 1.1e+02;
Conservative 0; Mismatches 0;
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JP 200250969-A/15.
synthetic construct
other sequences; artificial sequences.
1 (bases 1 to 20)
Keith, W.N.
                                               DNA

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/db_xref="taxon:32644"

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                                               20 bp
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BD225812
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                                                             Mammalian telomerase.
BD176152
BD176152.1 GI:29121856
JP 2002272489-A/11.
unidentified
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Best Local Similarity
                                                                                                                                                                inclassified.
                                                                                                                                                unidentified
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                                               BD176152
                                                          DEFINITION
ACCESSION
VERSION
KEYWORDS
SOURCE
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              RESULT 171
BD176152/c
LOCUS
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AUTHORS
TITLE
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VERSION
KEYWORDS
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TITLE
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PAT 07-SEP-2000
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                                                        PAT 12-JUN-2003
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Patent: WO 9938964-A 15 05-AUG-1999;
KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH (GB)
Location/Qualifiers
                                                                                                                                                  Unknow...
Unclassfiled.
1 (bases 1 to 20)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Wammalian telomerase
Mammalian telomerase
Patent: US 6548298-A 40 15-APR-2003;
Locaion/Qualifiers
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4.4%; Score 20; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 1.1e+02;
Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                    Length 20;
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/mol_type="unassigned DNA"
/moleare="taxon:32630"
/noTe="primer"
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synthetic construct
other sequences, artificial sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             other sequences; artificial sequences
                                                                                                                                                                                                                                                           /organism="unknown"
/mol_type="unassigned RNA"
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Sequence 45 from Patent WO9938964.
AX019591.1 GI:10043505
                                                        AR306488 20 bp Sequence 40 from patent US 6548298.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              AX019561 20 bp
Sequence 15 from Patent W09938964.
AX019561 GI:10043475
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                                                                                                                                                                                                                                                                                                                                                              41 TITGICTAACCCTAACTGAG 60
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                                                                                                  AR306488.1 GI:31696327
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synthetic construct
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Matches 20; Conserv
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                                                                                                                                Unknown.
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AX019591
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                                                                       DEFINITION
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                               RESULT 177
AR306488/c
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ORGANISM
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TITLE
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FEATURES
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KEYWORDS
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Unclassified.

1 (bases 1 to 20)

Willeponteau B., Feng, J., Funk, W. and Andrews, W.H. Mammalian telomerase
Patent: US 5583016-A 4 10-DEC-1996;

Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                  Unknow...
Unclassfiled.
1 (bases 1 to 20)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Mammalian telomerase
Patent: US 5583016-A 7 10-DEC-1996;
Location/Qualiflers
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Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H. Mammalian telomerase
Patent: US 6548298-A 7 15-APR-2003;
Location/Qualifiers
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100.0%; Pred. No. 1.1e+02;
tive 0; Mismatches 0; Indels
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4.4%; Score 20; DB 1; L.
Best Local Similarity 100.0%; Pred. No. 1.1e+02;
Matches 20; Conservative 0; Mismatches 0;
                                                                                                                                                 Score 20; DB 1; L
Pred. No. 1.1e+02;
                                                                                                                                              Similarity 100.0%; Pred. No. 1.1
0; Conservative 0; Mismatches
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/organism="unknown"
/mol_type="unassigned DNA"
                                                                                                        /organism="unknown"
/mol_type="unassigned DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 7 from patent US 6548298.
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Sequence 7 from patent US 5583016.
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/organism="unknown"
/mol_type="genomic DNA"
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Unclassified.
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Best Local Similarity
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ORGANISM
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AR306459/c
                                                                                                                                                                                                                                                                                                                       DEFINITION
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                                                              JOURNAL
FEATURES
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131754/c
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20 bp DNA linear PAT 27-AUG-2002
Method for detecting and inhibiting RNA component of telomerase.
BD023703
BP023703.1 GI:22564926
JP 2001507229-A/7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BD023702 20 Ap DNA linear PAT 27-AUG-2002 Method for detecting and inhibiting RNA component of telomerase. BD023702
WEINRICH
PC C12N15/09, A61K9/08, A61K31/7105, A61K45/00, A61K48/00, A61P35/00,
PC C12NS/10,
PC C12NS/10,
PC C12NS/10,
PC C12NS/12, C12Q1/68, C12Q1/68, C12N15/00, C12NS/00 CC
Strandedness: Single;
CC Topology: Linear;
CC Topology: Linear;
PC Note= 'oligo 14ab'
PH Key
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C12N5/10,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  unidentified unclassified.

unclassified.
1 (bases 1 to 20)
Kim, W., W., F., Kealey, J.T., Pruzan, R. and Weinrich, S.L. Kim, Nu, F., Kealey, J.T., Pruzan, R. and Weinrich, S.L. Method for detecting and inhibiting RNA component of telomerase Patent: JP 2001507229-A 6 05-JUN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PN JP 2001507229-A/6
PD 05-JUN-2001
PF 19-DEC-19997 JP 1998529003
PR 20-DEC-1996 US 08/770565 PI
NAM WOO KIM, FRED WU, JAMES T KEALEY, RONALD PRUZAN, SCOTT L PI
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PC C12N15/09,A61K9/08,A61K31/7105,A61K45/00,A61K4
PC C12N5/10,
PC C12N9/12,C12Q1/68,C12Q1/68,C12N15/00,C12N5/00
Strandedness: Single;
CC Topology: Linear;
CC Topology: Linear;
PC Fince Coligo 16ab'
PF KSP
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1.1e+02;
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4.4%; Score 20; DB 1; Lu
Best Local Similarity 100.0%; Pred. No. 1.1e+02;
Matches 20; Conservative 0; Mismatches 0;
                                                                                                                                    Location/Qualifiers.
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4.4%; Score 20; DB
Best Local Similarity 100.0%; Pred. No. 1.1
Matches 20; Conservative 0; Mismatches
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
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/organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
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JP 2001507229-A/6.
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JOURNAL
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      bD023699
      20 bp
      DNA
      linear
      PAT 27-AUG-20

      BD023699
      DNA
      component of telomerase.

      Keith, W.N.
Promoter regions of the mouse and human telomerase rna component
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 (bases 1 to 20)
Kim,N.W., Wu,F., Kealey,J.T., Pruzan,R. and Weinrich,S.L.
Method for detecting and inhibiting RNA component of telomerase
Patent: JP 2001507229-A 3 05-JUN-2001;
GERON CORP
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PD 05-JUN-2001
PP 19-DEC-1997 JP 1998529003
PR 20-DEC-1996 US 08/770564, 20-DEC-1996 US 08/770565 PI
NAM WOO KIM, FRED WU, JAMES T KEALEY, RONALD PRUZAN, SCOTT L PI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Au, J.L. and Wientjes, G.
Compositions active in telomere damage comprising a taxane and telomerase inhibitor
Patent: WO 0074667-A 6 14-DEC-2000;
Au, Jessie L.S. (US)
Location/Qualifiers
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                                      genes
Patent: WO 9938964-A 45 05-AUG-1999;
KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH (GB)
1. 20
//organism="synthetic construct"
//mol_type="unassigned DNA"
//do_type="unassigned DNA"
//do_xref="taxon:32630"
//note="Oligonucleotide"
                                                                                                                                                                                                                                    ch
    4.4%; Score 20; DB 1; Length 20;
1 Similarity 100.0%; Pred. No. 1.1e+02;
20; Conservative 0; Mismatches 0; Indels
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4.4%; Score 20; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 1.1e+02;
Matches 20; Conservative 0; Mismatches 0; Indels
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/organism="synthetic construct"
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
/note="primer/probe"
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synthetic construct
other sequences; artificial sequences.
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Sequence 6 from Patent WO0074667.
AXO58271
AXO58271.1 GI:12310770
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JP 2001507229-A/3.
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BD023699/c
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                                                                                                                                                                                                                                                                                                                                                    Zea mays
Zea mays
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PN JP 2001509681-A/11
PD 24-JUL-2001
PP 04-APR-1997 1998542718
PF SCOTT WEINFYLEH FEBWARD M ATKINSON III, SERGE P LICHTSTEINER, PI ALAIN P VASSEROT, RONALD A PRUZAN, JAMES T KEALEY PC C12N15/54, C12N9/12, C07K16/40, C12Q1/68, C07K14/47 CC Strandedness:
                                                                                                                                                                                                                                                                                                                                                                                                                              Spermatophyta; Magnoliophyta; Lillopsida; Poales; Poaceae; PACCAD clade; Panicoideae; Andropogoneae; Zea. Libases 1 to 20)
Weinrich, S.L., Lil, E.M.A., Lichtsteiner, S.P., Vasserot, A.P., Pruzan, R.A. and Kealey, J.T.
Puried telomerase
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                                                                                                                                                                                                                       linear
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GERON CORP
                                                                                                                                                                                                                       DNA
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/db_xref="taxon:4577"
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                                 159 TCTAGAGCAAACAAAAATG 178
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                                                                  20 TCTAGAGCAAACAAAAATG 1
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JP 2001509681-A/11.
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Purified telomerase.
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I (bases 1 to 20)

IS Kim, N w., wu, F. Kealey, J.T., Pruzan, R. and Weinrich, S.L.

Method for detecting and inhibiting RNA component of telomerase

Method for detecting and inhibiting RNA component of telomerase

D 2001507229-A 13 05-JUN-2001,

D 2001507229-A/13

PD 05-JUN-2001

PP 19-DEC-1997 JP 1998529003

PR 20-DEC-1996 US 08/770564,20-DEC-1996 US 08/770565 PI

NAM WOO KIM, FRED WU, JAMES T KEALEY, RONALD PRUZAN, SCOTT L PI

WEINRICH

PC C12NNJ/10, C12Q1/68, C12Q1/68, C12N15/00, C12N5/00 CC

Strandedness: Single;

CC Topology: Linear;

CC Topology: Linear;

CC Topology: Linear;

CC Topology: Linear;

CC Chote= 'Oilgo 20/21'

FH Key
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C12N5/10,
                        unidentified
unclassified.
1 (bases 1 to 20)
Kim,N.W., Wu,F., Kealey,J.T., Pruzan,R. and Weinrich,S.L.
Method for detecting and inhibiting RNA component of telomerase
Patent: JP 2001507229-A 7 05-JUN-2001;
GERON CORP
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PD 05-UTN-2001
PF 19-DEC-1997 UP 1998529003
PR 20-DEC-1996 US 08/770564, 20-DEC-1996 US 08/770565
NAM WOO KIM, FRED WU, JAMES T KEALEY, RONALD PRUZAN, SCOTT L PI
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PC C12N5/10,
PC C12N9/12,C12Q1/68,C12Q1/68,C12N15/00,C12N5/00 CC
Strandedness: Single;
CC Topology: Linear;
CC Topology: Linear;
FH Key Location/Qualifiers.
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4.4%; Score 20; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 1.1e+02;
Matches 20; Conservative 0; Mismatches 0; Indels
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1. 20
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/organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
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JP 2001507229-A/13.
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  unidentified
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BD023709/c
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BD023705/c
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BD071093
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JOURNAL
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TITLE
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                                                  linkages are replaced by N-(2-aminoethyl)glycine units linked
              Topology: Linear; /desc = 'peptide nucleic acid (PNA), where (deoxy(ribose- CC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 (bases 1 to 19)
Silverman, R.H., Kondo, S., Cowell, J.K., Li, G. and Torrence, P.F.
RNase L activators and antisense oligonucleotides effective to
treat telomerase-expressing malignancies
Patent: US 6468989-A 2 22-0CT-2002;
Location/Qualifiers
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Kim,N.Woo., Wu,F., Kealey,J.T., Pruzan,R. and Weinrich,S.L.

Methods for detecting the RNA component of telomerase

Patent: US 5846723-A 9 08-DEC-1998;

Location/Qualifiers
                                              C linkages are replaced by N-(2-aminoethyl)glycine
CC to
CC nucleotide bases via glycine amino N through a
methylenecarbonyl linker'
FH Key Location/Qualifiers
1. 20
FT source /organism='Unidentified'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 19; DB 1; Length 19;
Pred. No. 1.2e+02;
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Pred. No. 1.1e+02;
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100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                          Sequence 9 from patent US 5846723.

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/organism="unknown"
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Strandedness: Single;
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Best Local Similarity 100.03
Matches 19; Conservative
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Best Local Similarity 100.
Matches 20; Conservative
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AR241175/c
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Modulation of mammalian telomerase by peptide nucleic acids. BD071093 BD071093 GI:22616696 Unidentified unidentified
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                                                                                                                                                                                                                                                                                          BD023705 17-AUG-2
Method for detecting and inhibiting RNA component of telomerage.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              unclassified.

1 (bases 1 to 19)

Kim, N.W., Wu, F., Kealey, J.T., Pruzan, R. and Weinrich, S.L.
Method for detecting and inhibiting RNA component of telomerase
Patent: JP 2001507229-A 9 05-JUN-2001;
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PN JP 20012029-A/9
PN JP 20012029-A/9
PD 05-JUN-2001
PF 19-DEC-1997 JP 1998529003
PR 20-DEC-1996 US 08/770565
NAM WCO KIM, FRED WU, JAMES T KEALEY, RONALD PRUZAN, SCOTT L PI
WEINRICH
PC CL2N15/09, A61K9/08, A61K31/7105, A61K45/00, A61R97
PC CL2N15/09, A61K9/08, C12Q1/68, C12N15/00, C12N5/00 CC
Strandedness: Single;
CC Topology: Linear;
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Query Match 4.2%; Score 19; DB 1; Length 19; Best Local Similarity 100.0%; Pred. No. 1.2e+02; Matches 19; Conservative 0; Mismatches 0; Indels
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100.0%; Pred. No. 1.2e+02;
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4.2%; Score 19; DB
Best Local Similarity 100.0%; Pred. No. 1.2
Matches 19; Conservative 0; Mismatches

    .19
    /organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"

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JP 2001517929-A/59
09-0CT-2001
09-APR-1997 JP 1997536487
09-APR-1996 US 08/630019
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JP 2001507229-A/9.
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To (bases 1 to 20)
Weinrich, S.L., Atkinson, E.M. III, Lichtsteiner, S.P., Vasserot, A.P., Pruzan, R.A. and Kealey, J.T.
Purzán, R.A. and Kealey, J.T.
Purified telomerae.
Patent: US 5968506-A 14 19-OCT-1999;
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Spermatophyta, Magnoliophyta, Liliopsida, Poales, Poaceae, PACCAD
clade, Panicoideae, Andropogoneae, Zea.
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Weinrich, S.L., Iii, E.M.A., Lichtsteiner, S.P., Vasserot, A.P., Pruzan, R.A. and Kealey, J.T.
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4.2%; Score 19; DB 1; L
Best Local Similarity 100.0%; Pred. No. 1.3e+02;
Matches 19; Conservative 0; Mismatches 0;
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Sequence 14 from patent US 5968506.
AR079898
                                                                                                                                                                                                                        Patent: WO 9845450-A 7 15-OCT-1998;
GERON CORP (US)
                                                                                                                                                                                                                                                                                                                                       /mol_type="unassigned DNA"
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/mod_base=OTHER
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/wol_type="unassigned DNA"
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                                                                                                  unidentified
unclassified.
1 (bases 1 to 20)
Atkinson, B.M. and Kealey, J.T.
PURIFIED TELOMERASE
                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        361 AGGCCGCAGGAAGAGAAC 379
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JP 2001509681-A/7.
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Zea mays
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       A84597
A84597.1 GI:6733513
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Matches 19; Conserv
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AR079898/c
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BD058138/c
  ACCESSION
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KEYWORDS
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                                                         UdAMES C NURLUN
CO7K14/00,4618/16,C12Q1/68
Strandedness: Single;
Topology: Linear;
Modulation of mammalian telomerase by peptide nucleic acids FH
Location/Qualifiers
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synthe
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JERRY W SHAY, WOODRING E WRIGHT, MIECZYSLAW A PIATYSZEK, DAVID
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PR 21-APR-1997 US 60/044507,03-FEB-1998 US 09/018125
FOBERT H SILVERMAN, SELJI KONDO, JOHN K COWELL, GUIYING LI, PAUL
PI TORRENCE
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Description of Artificial Sequence: oligonucleotide
Location/Qualifiers
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    .19
    /organism="unidentified"

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/db_xref="taxon:32644"
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Sequence 7 from Patent WO9845450.
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                         COREY,
JAMES C NORTON
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Best Local Similarity
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Gaps

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Indels

Length 20;

PAT 31-AUG-2000

linear

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Gaps

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PAT 27-AUG-2002

linear

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Keith, W.N.
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AX019595
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BD225828/c
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                                                                      PN JP 2001509681-A/7
PN JP 2001509681-A/7
PD 24-UUD-2001
PF 04-APR-1997 JP 1998542718
PI SCOTT L WEINRICH, EDWARD M ATKINSON III, SERGE P LICHTSTEINER,
PI ALAIN P VASSEROT, ROWALD A PRUZAN, JAMES T KEALEY PC
CLIN15/54, C12N9/12, C07K16/40, C12Q1/68, C07K14/47 CC Strandedness:
Single,
CC Tropology: Linear;
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human telomerase RNA component genes.
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PC A61PS5/00, C12N1/15, C12N1/19, C12N1/21, C12NS/10, C12P21/02 PC A61PS5/00, C12N1/13, C12N1/19, C12N1/21, C12NS/10, C12P21/02 PC A61K35/76, A61K1:522), C12N1S/00, A61K37/02, C12NS/00 CC Description of Artificial Sequence:Oligonucleotide FH Key Location/Qualifiers
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4.2%; Score 18.8; DB 1; Length 22;
Best Local Similarity 90.9%; Pred. No. 1.5e+02;
Matches 20; Conservative 0; Mismatches 2; Indels
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100.0%; Pred. No. 1.3e+02;
tive 0; Mismatches 0;
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
Purified telomerase
Patent: JP 2001509681-A 7 24-JUL-2001;
GERON CORP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /mol_type="genomic DNA"
/db_xref="taxon:4577"
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/organism="Zea mays"
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                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
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JP 2002509699-A/49.
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synthetic construct
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Best Local Similarity
Matches 19; Conserv
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PF 29-JAN-1999 JP 200529424
PR 29-JAN-1998 GB 9801902.9
PI WILLIAM NICOL KEITH
PC 212N15/09, A61K31/711, A61K35/76, A61K38/00, A61K45/00, PC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       bp DNA linear PAT 17-JUL-200
human telomerase RNA component genes.
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CANCER RESEARCH CAMPAIGN TECHNOLOGY LTD
OS ATLIFICIAL Sequence
PN JP 2002509699-A/31
                                                                                                                                                                                                                                                                                                                                                  Promoter regions of the mouse and human telomerase rna component
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PC A61F35/00,C12N1/15,C12N1/19,C12N1/21,C12N5/10,C12P21/02 PC
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PC (A61K35/76,A61K31:522),C12N15/00,A61K37/02,C12N5/00 CC
Description of Artificial Sequence: Primer
FH Key Location/Qualifiers
FT source 1. .20
/organism='Artificial Sequence'.
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                                                                                                                                                                                                                                                                                                                                                                                           Patent: WO 9938964-A 49 05-AUG-1999;
KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH (GB)
Location/Qualifiers
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                                                                                                                                   linear
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/organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /organism="synthetic construct'
/mol_type="unassigned DNA"
/db_xref="taxon:12630"
/noTe="Oligonucleotide"
                                                                                                                                   DNA
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synthetic construct
synthetic construct
other sequences; artificial sequences.
1 (bases 1 to 20)
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synthetic construct
other sequences; artificial sequences.
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Sequence 49 from Patent WO9938964.
AX019595
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GGCCTGGGAGGGTAATGGCC 22
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Best Local Similarity 90.91
Matches 20; Conservative
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US09436060A-16.rge.sl

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/organism="synthetic construct"
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AX019580/c
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PF 29-JAN-1999 JP 2000529424

PR 29-JAN-1999 GB 9801902.9

PI WILLIAM NICOL KEITH

PC C12N15/09, A61K31/7105, A61K35/76, A61K38/00, A61K45/00, PC

A61F48/00,

PC A61F935/00, C12N1/15, C12N1/19, C12N1/21, C12N5/10, C12P21/02 PC

A61F35/76, A61K31:522), C12N15/00, A61K37/02, C12N5/00 CC

Description of Artificial Sequence: Primer

KeV Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                Promoter region of mouse and human telomerase RNA component genes Patent: JP 2002509699-A 34 02-ARR-2002;
CANCER RESEARCH CAMPAIGN TECHNOLOGY LTD
OS Artificial Sequence
PN JP 2002509699-A/34
PD 02-ARR-2002
PP 29-JAN-1999 JP 2000529424
PR 29-JAN-1999 GB 9801902.9
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Promoter region of mouse and human telomerase RNA component genes.
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Patent: WO 9938964-A 31 05-AUG-1999;
KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH (GB)
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4.1%; Score 18.4; DB 1; Length 20; 95.0%; Pred. No. 1.4e+02; tive 0; Mismatches 1; Indels
Score 18.4; DB 1; Length 20;
Pred. No. 1.4e+02;
0; Mismatches 1; Indels
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/organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
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synthetic construct
other sequences; artificial sequences.
(bases 1 to 20)
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Sequence 31 from Patent WO9938964.
AX019577
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                                                           102 TTCTCGCTGACTTTCAGCGG 121
20 TTCTCGCTGACTTCCAGCGG 1
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    4.18;
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synthetic construct
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Best Local Similarity 95.0%
Query Match
Best Local Similarity 95.0°
Matches 19; Conservative
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LOCUS
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AX019577/c
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BD196339 18-bp DNA linear PAT 17-JUL-2003 Vertebrate telomerase genes and proteins and uses thereof. BD196339
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Description of Artificial Sequence:Synthesized Amplification
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synthetic construct
synthetic construct
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construct
synthetic and Bowtell, D.
Synthetic and Bowtell, D.
Synthetic and Bowtell, D.
Artificial Sequence
Natificial Sequence
Solossalva
Colossalva

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KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH (GB)
Location/Qualifiers
                                                                                                                                                                      Length 20;
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                                                                                                                                                                 Query Match
4.1%; Score 18.4; DB 1;
Best Local Similarity 95.0%; Pred. No. 1.40+02;
Matches 19; Conservative 0; Mismatches 1;
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4.1%; Score 18.4; DB 1;
Best Local Similarity 95.0%; Pred. No. 1.46+02;
Matches 19; Conservative 0; Mismatches 1;
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/organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="taxon:32630"
/note="primer"
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
/note="primer"
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Sequence 34 from Patent WO9938964.
AX019580
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20 TTCTCGCTGACTTCCAGCGG 1
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JP 2002514928-A/73.
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C12N15/09,A61K31/7105,A61K31/711,A61K35/76,A61K38/00,A61K45/00, PC
                                                                                                                                                        Promoter region of mouse and human telomerase RNA component genes Patent: 4P 200220599-A 47 02-APR-2002;
CANCER RESEARCH CAMPAIGN TECHNOLOGY LTD
OS Artificial Sequence
     Promoter region of mouse and human telomerase RNA component genes
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PC A61P35/00, C12N1/15, C12N1/19, C12N1/21, C12NS/10, C12P21/02 PC A61P35/00, C12N1/15, C12N1/19, C12N1/21, C12NS/10, C12P21/02 PC C1201/68//C12N9/12, C12N15/00, A61K37/02, C12N5/00 CC Description of Artificial Sequence:Oligonucleotide FH Key Location/Qualifiers
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KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH (GB)
Location/Qualifiers
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86.4%; Pred. No. 1.9e+02;
tive 0; Mismatches 3;

    .22
    ^organism="synthetic construct"
/mol_type="genomic DNA"
    /db_xref="taxon:32630"

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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
/noTe="Oligonucleotide"
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1 (bases 1 to 22)
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Sequence 47 from Patent W09938964.
AX019593
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02-APR-2002
29-JAN-1999 JP 2000529424
29-JAN-1998 GB 9801902.9
WILLIAM NICOL KEITH
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                                       BD225844.1 GI:33035614 JP 2002509699-A/47.
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Best Local Similarity 86.4°
Matches 19; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unidentified
JP 2001517929-A/9
09-0CT-2001
09-0AR-1997 JP 1997536487
09-ARR-1996 US 08/630019
JERRY W SHAY, WOODRING E WRIGHT, MIECZYSLAW A PIATYSZEK, DAVID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          unidentified
unclassified.
1 (bases 1 to 18)
Shay,J.W., Wright,W.E., Piatyszek,M.A., Corey,D. and Norton,J.C.
Modulation of mammalian telomerase by peptide nucleic acids
Patent: JP 2001517929-A 9 09-OCT-2001;
Primer Design
based on EST Sequence Genbank Accession Number AA281296 FH
Location/Qualifiers
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CC nucleotide bases via glycine amino N through a
methylenecarbonyl linker'
FH Key Location/Qualifiers
FT source location/Qualifiers
FT source /organism='Unidentified'.
                                                                         'organism='Artificial Sequence'
                                                                                                                                                                                          4.0%; Score 18; DB 1; Length 18;
100.0%; Pred. No. 1.4e+02;
Live 0; Mismatches 0; Indels
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Pred. No. 1.4e+02;
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    18
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        /db_xref="taxon:32630"

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100.0%; Pred. No....
0; Mismatches

    .18
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Strandedness: Single;
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JP 2001517929-A/9.
unidentified
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Best Local Similarity 100.09
Matches 18; Conservative
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JAMES C NORTON
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Best Local Similarity 100.0
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BD071043/c
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C12Q1/48,C12Q1/68,G01N33/15,G01N33/48,G01N33/50//C07K14/47, PC
C07K16/40,
C12N1/19,C12N1/21,C12N5/10,C12N9/12,C12P21/08,(C12N1/19, PC
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                   PAT 18-JUN-2001
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C12N15/09,A61K31/70,A61K38/55,A61K39/395,A61K39/395,A61K48/00,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LCKL:84), LCL2NS/10,CL2N9/12,CL2P21/08,(CL2N1/19, PC CC CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,CL2R1:19),(CL2N9/12,C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        08/844.419, PR
                                                                                                                                                           unclassified.

Unclassified.

I (bases 1 to 18)

SS Calvin, B.H. and William, H.A.

Human telomerase catalytic subunit promoter

Human telomerases catalytic subunit promoter

B thuman telomerases catalytic subunit promoter

IAL DERON CORP, UNIVERSITY TECHNOLOGY CORP

OS Unidentified

PN JP 1999253177-A/201

PD 21-SEP-1999

PN JP 1999253177-A/201

PP 15-OCT-1998 JP 1998320169

PR 01-OCT-1998 JP 1998320169

PR 01-OCT-1998 JP 199846.017,06-MAY-1997 US 08/851.843, PR 09-MAY-1997 US 08/866.017,06-MAY-1997 US 08/911.312, PR 14-AUG-1997 US 08/9112.911,14-AUG-1997 US 08/9113.12, PR PR SECHI, JOCHIMU RINGNER, TORU NAKAMURA, KAREN B CHAPMAN, PI GREG
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Morin,G.B. and Andrews,W.H.
Promoter for telomerase reverse transcriptase
Patent: US 6610839-A 543 26-AUG-2003;
Location/Qualifiers
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Location/Qualifiers
                   E36993 18 bp DNA l
Human telomerase catalytic subunit promoter.
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Pred. No. 1.7e+02;
0; Mismatches 1;
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Sequence 543 from patent US 6610839.
AR390670.1 GI:40112602
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                                                                                            E36993.1 GI:13022956
UP 1999253177-A/201.
unidentified
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Best Local Similarity 94.4%;
Matches 17; Conservative
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                                   DEFINITION
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AR390670/c
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E36993/c
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C12N5/10,
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Kim, W., W., W., W., W. K.

Kim, N., W., W., W., K.

Kim, N., W., W., W.

Method for detecting and inhibiting RNA component of telomerase patent: JP 2001507229-A 14 05-JUN-2001;

GERON CORP

PN JP 2001507229-A/14

PD 05-JUN-2001

PP 19-DEC-1997 JP 1998529003

PR 20-DEC-1996 US 08/770564, 20-DEC-1996 US 08/770565 PI

NAM WOO KIM, FRED WU, JAMES T KEALEY, RONALD PRUZAN, SCOTT L PI
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                                                                                                                                                                                                                             Unknown.
Unclassified.
Unclassified.
Kim,N.Woo., Wu,F., Kealey,J.T., Pruzan,R. and Weinrich,S.L.
Kim,N.Woo., Wu,F., Kealey,J.T., Pruzan,R. and Weinrich,S.L.
Methods for detecting the RNA component of telomerase
Patent: US 5846723-A 14 08-DEC-1998;
Location/Qualifiers
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PC C12N5/10,
C C12N5/12,C12Q1/68,C12Q1/68,C12N15/00,C12N5/00 CC
Strandedness: Single;
CC Topology: Linear;
CC Topology: Linear;
FH Key Location/Qualifiers.
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larity 100.0%; Pred. No. 1.5e+02;
Conservative 0; Mismatches 0;
                                                                               DNA
                                                                    Sequence 14 from patent US 5846723.
AR063838
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JP 2001507229-A/14.
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                           RESULT 205
AR063838/c
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THOMAS GREG B

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פטנים בחשבים PAT 17-JUL-2003
Vertebrate telomerase genes and proteins and uses thereof.
BD196340
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PC A61K38/00, A61K31/7088, A61K39/00, A61P35/00, A61P43/00,
PC C07K5/10,
PC C07K5/10,
PC C07K5/10,
PC C12K15/09,
PC C12Q1/02, C12Q1/48, C12Q1/68, G01N33/15, G01N33/50, G01N33/53, PC G01N33/53,
          PAT 31-JAN-2002
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BD196340.1 GI:33006110
JP 2002514928-A/74.
Synthetic construct
SM synthetic construct
cocher sequences; artificial sequences.
Chases 1 to 21)
S Kilian,A. and Bowtell,D.
Vertebrate telomerase genes and proteins and uses thereof
D 2002514928-A 74 21-MAY-2002;
CAMBIA BIOSYSTEWS LLC,PETER MACCALLUM CANCER INSTITUTE
OS Artificial Sequence
FN JP 2002514928-A/74
FN JP 2002514928-A/74
FN JP 2002514928-A/74
FN JP 2002514928-B/74
FN JP 2002514938-B/74
FN JP 2002514938-B/74
FN JP 2002514938-B/74
FN JP 2002514938-B/74
FN JP 200251498-B/74
FN JP 200251498-B/74
FN JP 200251498-B/74
FN JP 200251498-B/74
FN JP 200251499-B/74
FN JP 2002514
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CC Topology: Linear;
FH Key Location/Qualifiers
FT source L. 18
FT /organism='Unidentified'.
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             linear
Human telomerase catalytic subunit.
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/db_xref="taxon:32644"
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                                                                                                      BD011244.1 GI:18639617
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Best Local Similarity 94.4
Matches 17; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 (bases 1 to 18)
Cech,T.R., Lingmer,J., Nakamura,T., Chapman,K.B., Morin,G.B.,
Harley,C.B. and Andrews,W.H.
Cells immortalized with telomerase reverse transcriptase for use in
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Human telomerase catalytic subunit
Patent: EP 1333094-A 53 06-AUG-2003;
Geron Corporation (US); University Technology Corporation (US)
Location/Qualifiers
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3.6%; Score 16.4; DB 1; Length 18;
Best Local Similarity 94.4%; Pred. No. 1.7e+02;
Matches 17; Conservative 0; Mismatches 1; Indels
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Patent: US 6617110-A 543 09-SEP-2003;
Location/Qualiflers
                                                                                                                                                                                                                                                                                                                                                                                            Sequence 543 from patent US 6617110. AR393284
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/mol_type="unassigned DNA"
/db xref="taxon:32644"
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Sequence 543 from Patent EP1333094.
AX810578
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/organism="unknown"
/mol_type="genomic DNA"
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Matches 17; Conservative (
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AX810578/c
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BD011244/c
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THOMAS GREG B

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bp DNA linear PAT 17-JUL-2003
human telomerase RNA component genes.
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PN 19 20255099-A/17
PD 02-APR-2002
PF 29-JAN-1999 JP 2000529424
PR 29-JAN-1998 GB 9801902.9
PI WILLIAM NICOL KEITH
PC 212N15/09,A61K31/7105,A61K31/711,A61K35/76,A61K38/00,A61K45/00, PC
                                                                                                                                                                                        linkages are replaced by N-(2-aminoethyl)glycine units linked
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Parameter region of mouse and human telomerase RNA component genes Parent: JP 20025099-A 17 02-APR-2002;
CANCER RESPARCH CAMPAIGN TECHNOLOGY LTD
OS Artificial Sequence
PN JP 2002509699-A/17
PP 29-JAN-1999 JP 2000529424
PR 29-JAN-1999 GB 9801902.9
                                                                                                                      Strandedness: Single;
Topology: Linear;
/desc = 'peptide nucleic acid (PNA), where (deoxy(ribose- CC
phosphate
              09-APR-1997 JP 1997536487
09-APR-1996 US 08/630019
JERRY W SHAY, WOODRING E WRIGHT, MIECZYSLAW A PIATYSZEK, DAVID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A61K49/00,
oc A61P35/00,C12N1/15,C12N1/19,C12N1/21,C12N5/10,C12P21/02 PC
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PC (A61K35/76,A61K31:522),C12N15/00,A61K37/02,C12N5/00
Description of Artificial Sequence: Primer
FH Key Location/Qualifiers
FT source 1. 20
/ Organism='Artificial Sequence'.
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CC nucleotide bases via glycine amino N through a
methylenecarbonyl linker'
FH Key l. 16
FT source l. 16
/organism='Unidentified'.
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         other sequences; artificial sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; Prec. .v..
                                                                                    JAMES C NORTON
C07K14/00, A61K38/16, C12Q1/68
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/organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
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and
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synthetic construct
synthetic construct
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 DEFINITION
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Gorman, C.M. and Groskreutz, D.J.
Non-endocrine animal host cells capable of expressing variant proinsulin and processing the same to form active, mature insulin and methods of culturing such cells
Patent: US 6348327-A 13 19-FBB-2002;
Location/Qualifiers
                    ANDRZEJ KILIAN, DAVID BOWTELL
CI2NI5/54, CI2N9/12, A61K38/45, C07K16/40, C12Q1/68, C12Q1/48, PC
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Shay, "W. Wright, W.E., Piatyszek, M.A., Corey, D. and Norton, J.C. Modulation of mammalian telomerase by peptide nucleic acids Patent: JP 2001517929-A 26 09-OCT-2001, GERON CORP OS Unidentified
                                                                                      Description of Artificial Sequence:Synthesized Amplification
                                                                                                         Primer Design
based on EST Sequence GenBank Accession Number AA281296 FH
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Modulation of mammalian telomerase by peptide nucleic acids
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                                                                                                                                                                         /organism='Artificial Sequence' Location/Qualifiers
                                                                                                                                                                                                                                                                                                     Score 16.2; DB 1; Length 21; Pred. No. 2.1e+02;
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                                                                                                                                                                                                                                                                                                                                        3; Indels
                                                                                                                                                                                                                1. .21
/organism="synthetic construct"
/mol_type="genomic DNA"
/db_kref="taxon:32630"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA
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/organism="unknown"
/mol_type="unassigned DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21 bp Sequence 13 from patent US 6346327.
                                                                                                                                            Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                        0; Mismatches
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   60/058287
                                                                                                                                                                                                                                                                                                                                                                           431 CAGGACTCGCCTCACACATGC 451
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JP 2001517929-A/26.
unidentified
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                                                                                                                                                                                                                                                                                                         3.6%;
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Best Local Similarity 85.7
Matches 18; Conservative
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Best Local Similarity 85.7
Matches 18; Conservative
09-SEP-1997 US
PI ANDEZEJ KII
PC CL2N15/54,C
CL2N15/11,
PC A61K31/70
CC Description
CC Dased on E9
CC based on E7
KEY
FT source
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Bougueleret, L.
Nucleic acid encoding a retinoblastoma binding protein (RBP-7) and polymorphic markers associated with said nucleic acid Patent: US 6399373-A 76 04-JUN-2002;
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Li.X.-J. and Li.S.-H.
Hybridoma cell line and monoclonal antibody for huntingtin protein
Patent: US 6355481-A 11 12-MAR-2002;
                                                             Gape
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85.0%; Pred. No. 2.3e+02;
tive 0; Mismatches 3; Indels
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                              Length 20
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                           Query Match
3.4%; Score 15.4; DB 1;
Best Local Similarity 94.1%; Pred. No. 2.2e+02;
Matches 16; Conservative 0; Mismatches 1;
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Sequence 76 from patent US 6399373,
AR211866
AR211866.1 GI:21515301
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AR199735
AR199735.1 GI:20249809
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Sequence 11 from Patent WO0078813.
AX060355
/db_xref="taxon:9606"
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Best Local Similarity 94.1.
Best Local 6; Conservative
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Best Local Similarity 85.0
Matches 17; Conservative
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AX060355
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20 bp DNA linear PAT 17-JUL-2003
Nucleic acid encoding retinoblastoma-binding protein (RBP-7) and
polymorphic marker relating to the nucleic acid.
BD221936
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JP 2002519027-A/75.
HOMO sapiens (human)
HOMO sapiens
Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
(homo sapiens)
Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
(homoalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Bougueleret, L.
Nucleic acid encoding retinoblastoma-binding protein (RBP-7) and
polymorphic marker relating to the nucleic acid
Patent: JP 2002519027-A 75 02-JUL-2002;
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                                                                                                                                                                                                                                                                                          Promoter regions of the mouse and human telomerase rna component
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PC C12N15/09.A01K67/027,C07K14/47,C07K16/18,C12N5/PC C1201/68,
PC G01N33/53,G01N33/566,C12N15/00,C12N5/00,C12N5/00,C12N15/00 CC Upstream amplification primer for SEQ 34, SEQ 55, SEQ 35, SEQ
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KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH (GB)
Location/Qualifiers
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                                                                                                                             linear
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/organism="synthetic construct"
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100.0%; Pred. No. 2e+02;
tive 0; Mismatches 0
                                                                                                                             DNA
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Sequence 17 from Patent WO9938964.
AX019563
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JP 2002519027-A/75
02-JUL-2002
30-JUN-1999 JP 2000557360
30-JUN-1998 US 60/091315
IE BOUGUELERET
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Location/Qualifiers
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                CTCGGCTCACACATGC 451
                                   CTCGGCTCACACATGC 16
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synthetic construct
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BD071036 15 bp DNA linear PAT 27-AUG-2002 Modilation of mammalian telomerase by peptide nucleic acids.
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PC C12N15/09, A61K9/08, A61K31/7105, A61K45/00, A61K48/00, A61P35/00,
PC C12N15/09, A61K9/08, C12Q1/68, C12N15/00, C12N5/00, CC
Strandedness: Single;
CC Topology: Linear;
CC Topology: Linear;
CC Topology: Linear;
PH Key
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JP 2001s17929-A/2
JP 2001s17929-A/2
09-CCT-2001
09-APR-1996 US 08/630019
JERRY W SHAY, WOODRING E WRIGHT, MIECZYSLAW A PIATYSZEK, DAVID
COREY,
JAMES C NORTON
COTK14/00, A61K38/16, C12Q1/68
Strandedness: Single,
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/desc = 'peptide nucleic acid (PNA), where (deoxy(ribose- CC
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unidentified
unidentified
unclassified.
1 (Dases 1 to 15)
Shay,J.W., Wright,W.E., Piatyszek,M.A., Corey,D. and Norton,J.C.
Patent: JP 2001517929-A 2 09-OCT-2001;
GERON CORP
       PR 20-DEC-1996 US 08/770564,20-DEC-1996 US 08/770565 PI NAM WOO KIM, FRED WU, JAMES T KEALEY, RONALD PRUZAN, SCOTT L PI
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FR Key Location/Qualifiers 1.15F
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Location/Qualifiers
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JP 2001517929-A/2.
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JP S001507259-A/10.
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                                                                                                                  Huntington disease cellular model: stably transfected pc12 cells expressing mutant huntingtin Patent: WO 0079813-A 11 28-DEC-2000; Emory University (US) Location/Qualifiers
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Kim.N.W., Wu.F., Kealey, J.T., Pruzan, R. and Weinrich, S.L.

Method for detecting and inhibiting RNA component of telomerase Patent: JP 2001507229-A 10 05-JUN-2001;

PN JP 2001507229-A/10

PD 05-JUN-2001

PF 19-DPG 2001
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/note="PCR primer"
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                                                                        other sequences; artificial sequences
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Sequence 10 from patent US 5846723,
AR063834
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/organism="unknown"
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   AX060355.1 GI:12405842
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synthetic construct
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Unclassified.
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AR063834/c
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As beauth a color of mammalian telomerase by peptide nucleic acids Modulation of mammalian telomerase by peptide nucleic acids patent; JP 2001517929-A 44 09-OCT-2001; GENON CORP OS Unidentified by D 09-OCT-2001; D 09-OCT-2001 by D 09-OCT-2001 b
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                                                                                                 Topology: Linear; //desc = 'poptide nucleic acid (PNA), where (deoxy(ribose- CC phosphate phosphate linkages are replaced by N-(2-aminoethyl)glycine units linked
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gape
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BD071078
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PI COREY,
PI JAMES C NORTON
PC COTAIA/00, AGIN38/16, C12Q1/68
CC Strandedness: Single,
CC Topology: Linear;
CC /desc = 'peptide nucleic acid (PNA), where (deoxy(pescare))
CC linkages are replaced by N-(2-aminoethyl)glycine ucc
CC nucleotide bases via glycine amino N through a CC methylenecarbonyl linker'
FH Key Location/Qualifiers
FT source / Organism='Unidentified'.
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Location/Qualifiers
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/db_xref="taxon:32644"

    .15
    /organism="unidentified"
    /mol_type="genomic DNA"
    /db_xref="taxon:32644"

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1 (bases 1 to 15)
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Best Local Similarity
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BD071078/c
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JP 2001517929-A/27.  
unidentified
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JAMES C NORTON
CO7K14/00,A61K38/16,C12Q1/68
Strandedness: Single;
Topology: Linear;
/desc = 'Peride nucleic acid (FNA), where (deoxy(ribose- CC
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JP 2001517929-A/27
09-OCT-2001
09-APR-1997 JP 1997536487
09-APR-1996 US 08/630019
JERRY W SHAY,WOODRING E WRIGHT,MIECZYSLAW A PIATYSZEK,DAVID
                                                                                                                                                             DD0/1039.1 GI:22616642
JP 2001517929-A/5.
unidentified
unidantified
unclassified.
1 (bases 1 to 15)
Shay, J. W., Wright, W.E., Piatyszek, M.A., Corey, D. and Norton, J.C.
Modulation of mammalian telomerase by peptide nucleic acids
Patent: JP 2001517929-A 5 09-0CT-2001;
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JERRY W SHAY,WOODRING E WRIGHT,MIECZYSLAW A PIATYSZEK,DAVID
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unclassified.
1 (bases 1 to 15)
Shay,J.W., Wright,W.E., Piatyszek,M.A., Corey,D. and Norton,J.C.
Modulation of mammalian telomerase by peptide nucleic acids
Patent: JP 2001517929-A 27 09-OCT-2001;
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Modulation of mammalian telomerase by peptide nucleic acids.
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CC CC CC CC nucleotide bases via glycine amino N through a methylenecarbonyl linker'
FR Key Location/Qualifiers
1.15
/ Organism='Unidentified'.
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    /organism='Unidentified'

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100.0%; Pred. No. 1.8e+02;
tive 0; Mismatches 0;
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JP 2001517929-A/S
09-0CT-2001
09-APR-1997 JP 1997536487
09-APR-1996 US 08/630019
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SYN 26-MAR-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1. .19
/organism="synthetic construct"
/mol type="other DNA"
/db_xref="taxon:32630"
/db_xref="forward primer for Japanese flounder microsatellite sequence Poli104MHFS"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PAT 17-JUL-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Fuji,K., Kobayashi,K., Mizuta,A., Hasegawa,O., Tabata,K., Sakamoto,T. and Okamoto,N. A genetic linkage map of the Japanease Flounder, (Paralichthys olivaceus)
Unpublished
2 (bases 1 to 19)
Mizuta,A., Tabata,K., Kobayashi,K., Fuji,K., Sakamoto,T. and
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Synthetic construct DNA, forward primer for Japanese flounder
microsatellite sequence Poli104HFS-M.
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3.3%; Score 14.8; DB 1;
Best Local Similarity 88.9%; Pred. No. 2.3e+02;
Matches 16; Conservative 0; Mismatches 2;
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synthetic construct
other sequences; artificial sequences.
1 (bases 1 to 18)
Kusu, M., Qiu,G. and Hunfrees,R.
acid differentiation
Patent: US 5688643-A 22 18-NOV-1997;
Location/Qualifiers
                                                                      /organism="unknown"
/mol_type="unassigned DNA"
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synthetic construct
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Best Local Similarity
Matches 16; Conserv
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AB175197
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                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unknown.
Unclassified.
1 (bases 1 to 18)
Schafer,M.P. and Reid,T.M.
Rapid and sensitive method for detecting histoplasma capsulatum
Rapid and sensitive method for detecting histoplasma capsulatum
Patent: US 646156-4 4 22-00T-2002;
Location/Qualifiers
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Oka,T., Matsunaga,H. and Yamane,A.
Method of nucleic acid-differentiation and assay kit for nucleic
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                                                 linear
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3.3%; Score 15; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 2.3e+02;
Matches 15; Conservative 0; Mismatches 0;
                                            AX613673 20 bp DNA
Sequence 4698 from Patent W002072882.
AX613673
                                                                                                                                                                                                                             Coronary chip
Patent: WO 02072882-A 4698 19-SEP-2002;
OGHAM GmbH (DE)
                                                                                                                                                                                                                                                                                                    1. .20
/organism="Homo sapiens"
/mol_type="unassigned DNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 4 from patent US 6469156.
AR241369
AR241369.1 GI:27287033
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Sequence 22 from patent US 5688643.
174343
174343.1 GI:3010484
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                                                                                            AX613673.1 GI:28409102
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                                                                                                                               Homo sapiens (human)
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Best Local Similarity 88.99
Matches 16; Conservative
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Unclassified.
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AR241369/c
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Voss,J. and Timm,J.
Methods and compositions for the screening of cell cycle modulators
Patent: WO 0120034-A 17 22-MAR-2001;
BASF AKTIENGESELLSCHAFT (DE)
Location/Qualifiers
   INNOGENETICS N.V. (BE) ; Enterprise Ireland (trading as Bioresearch
Ireland) (IE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17 bp DNA linear PAT 17-JUL-2003 on character imprinted by parent.
                                                                                                                                                                                                                                                                                                                                            PAT 02-APR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mus musculus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CARINE
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Andersson,L., Georges,M., Spincemaille,G. and Nezer,C.D.A. Selection of animal based on character imprinted by parent Patent: JP 2005335963-A 130 29-OCT-2002; UNIVERSITY OF LIEGE,MELICA HB, SEGHERS GENTEC NV
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JP 2002535963-A/130
29-CCT-2002
16-DEC-1999 JP 2000588390
16-DEC-1998 EP 98204291.3
LEIF ANDERSSON, MICHEL GEORGES, GEERT SPINCEMAILLE, PI
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                                                                                                                                              Length 17;
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100.0%; Pred. No. 2.4e+02;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                 0; Indela
                                                                      /organism="Aspergillus versicolor"
/mol_type="unassigned DNA"
/db_xref="taxon:46472"
                                                                                                                                              Query Match
3.1%; Score 14; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 2.4e+02;
Matches 14; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                              DNA
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/mol_type="unassigned DNA"
/db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                                                                       Sequence 17 from Patent W00120034. AX099957
                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                     Mus musculus (house mouse)
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Selection of animal based
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BD251610.1 GI:33061380
JP 2002535963-A/130.
Sus scrofa (pig)
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PC C12N15/09,A01K67
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Best Local Similarity 100.0%
Matches 14; Conservative
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AX099957/c
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Patent: JP 2002531582-A 23 24-SEP-2002;

ANTIGEN EXPRESS INC
OS Artificial Sequence
PN JP 2002531582-A/23
PD 24-SEP-2002
PP 24-SEP-2002
PP 24-SEP-2009
PP 04-DEC-1998 US 09/205995
PI minzhen kusu,gang qiu,robert hunfrees
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Aspergillus versicolor
Eukaryota; Fungi; Ascomycota; Pezizomycotina; Eurotiomycetes;
Eurotiales; Trichocomaceae; mitosporic Trichocomaceae; Aspergillus.
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Xu,M., Qiu,G. and Humphreys,R.
MHC class II antigen presenting cells containing oligonucleotides
WHich inhibit II protein expression
Patent: US 6368955-A 24 09-APR-2002;
Location/Qualifiers
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Weide,M.
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                                                                                                                                                             corresponding to a specific region of the mouse Ii gene. Key Location/Qualifiers Location/Qualifiers
                                                                                                                                                                                                                                     1. .18
/organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
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Patent: WO 0073499-A 21 07-DEC-2000;
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Sequence 24 from patent US 6368855.
AR205264
AR205264.1 GI:21502804
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/organism="unknown"
/mol_type="unassigned DNA"
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AX055663
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Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Butheria, Primates, Catarrhini, Hominidae, Homo.
                                                                                                                                                                                Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Cetartiodactyla, Suina, Suidae, Sus.
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                                                                                                                                                                                                                         Georges, M., Spincemaille, G. and Andersson, L. selecting animals for parentally imprinted traits batent: WO 0036143-A 130 22-JUN-2000; SEGHENSCENTEC N (BB); GEORGES MICHEL (BB); UNIV LIEGE SPINCEMAILLE GERRT (BB); MELICA HB (SE); ANDERSSON LEIF LOCATION/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Elis,J.H.
Method and reagent for the inhibition of grid
Patent: WO 0162911-A 129 30-AUG-2001;
RIBOZYME PHARMACEUTICALS, INC. (US); GLAXO GROUP LIMITED
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/organism="Sus scrofa"
/mol_type="unassigned DNA"
/moletype="taxon:9823"
/noFe="Polymorphism Tyrosine Hydroxylase gene"
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Pred. No. 2.4e+02;
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AX272560.1 GI:16545297
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Sequence 330 from Patent WO0162911.
AX272761
AX272761.1 GI:16545498
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/mol_type="unassigned RNA"
/db_xref="taxon:9606"
                                                                              Sequence 130 from Patent WO0036143.
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AX028311.1 GI:10189199
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Best Local Similarity 88.2%;
Matches 15; Conservative
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                                                                                                                                                                  Sus scrofa
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Unclassified.
1 (bases 1 to 17)
Zwick,M.G., Edington,B.E., McSwiggen,J.A., Merlo,P.Ann.Owens.,
Guo,L., Skokut,T.A., Young,S.A., Folkerts,O. and Merlo,D.J.
Nucleic acid encoding delta-9 desaturase
Patent: US 6350934-A 826 26-FEB-2002;
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                         /organism='Sus scrofa (pig)'.
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Sequence 826 from patent US 6350934.
AR196361
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PNA-DNA-PNA chimeric macromolecules
Patent: US 5700922-A 2 23-DEC-1997;
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /organism="unknown"
/mol_type="unassigned DNA"
                                                                                                                                                                                                                                                                                                                              17 bp
Sequence 2 from patent US 5700922.
186370

    .17
    /organism="Sus scrofa"
    /mol_type="genomic DNA"
    /db_xref="taxon:9823"

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                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Randi, A. M.
Randi, A. M.
Method and reagent for the inhibition of erg
Patent: WO OBB124-A 1949 22-NOV-2001,
RIBOZYME PHRMACEUTICALS, INC. (US); GLAXO GROUP LIMITED (GB)
Location/Qualifiers
                                                                                         Jarvis, T., von Carlowitz, I., Mcswiggen, J.A., Hamblin, P.A. and Ellis, J.H.
Method and reagent for the inhibition of grid
Patent: WO 0162911-A 330 30-AUG-2001;
RIBOZYME PHARMACEUTICALS, INC. (US); GLAXO GROUP LIMITED (GB)
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other sequences; artificial sequences.
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Sequence 1949 from Patent WO0188124.
AX423613
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Pna-dha-pna chimeric macromolecules
Patent: EP 1201676-A 2 02-MAY-2002;
PSISS PHARMACEUTICALS, INC. (US)
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/db_xref="taxon:9606"
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/db_xref="taxon:9606"
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/organism="Homo sapiens"
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Homo sapiens
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AX423613/c
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Shannon, M., Gu, Y. and Nguyen, C.T. Four human zinc-finger-containing proteins : mdz3, mdz4, mdz7 and
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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                                                                                                                                                      Length 17;
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1..17
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/db xref="taxon:32630"
/note="PNA analogue"
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Molecular Engines Laboratories (FR)
Location/Qualifiers
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Sequence 5862 from Patent WO03025176.
AX728175
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Aeomica, Inc. (US)
Location/Qualifiers
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Sequence 814 from Patent EP1281758.
AX688082
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/db_xref="taxon:9606"
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Mus musculus
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US09436060A-16.rge.sl

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PAT 07-SEP-2000
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Guida,M. and Hall,J.
Guida,M. and Guida,M. and Guida,M. and Hall,J.
Genetic typing of the human cytochrome P450 2A6 gene and related materials and methods
Patent: US 6492115-A 11 10-DEC-2002,
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                   Promoter regions of the mouse and human telomerase rna component
                                                             Gaps
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Patent: WO 9938964-A 40 05-AUG-1999;
KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH (GB)
Location/Qualifiers
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                             Length 38;
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                                                             Indels
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                            Query Match 3.1%; Score 13.8; DB 1; Best Local Similarity 72.0%; Pred. No. 3.2e+02; Matches 18; Conservative 0; Mismatches 7;
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Pred. No. 3.2e+02;
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/noTe="Oligonucleotide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA
                                                                                                                                                                                                                 DNA
                                                                                                                                                                                                                                                                                                                                   other sequences; artificial sequences
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Best Local Similarity 72.0%; Pred. No. 3.2e
Matches 18; Conservative 0; Mismatches
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AR264866
AR264866.1 GI:29693235
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Sequence 40 from Patent WO9938964.
AX019586
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/mol_type="genomic DNA"
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synthetic construct
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Matches 14; Conservative
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Unclassified.
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AX019586/c
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AR264867
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TITLE
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C12N15/09,A61K31/7105,A61K31/711,A61K35/76,A61K38/00,A61K45/00,PC
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Promoter region of mouse and human telomerase RNA component genes.
BD225837
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CANCER RESEARCH CAMPAIGN TECHNOLOGY LTD
OS Artificial Sequence
PD 02-ARF-2002
PD 02-ARF-2002
PR 29-JAN-1999 JP 2000529424
PR 29-JAN-1999 GB 9801902.9
PI WILLIAM NICOL KEITH
                                                                                                                                                                                                                                                                             Unknown.
Unclassified.
I (bases I to 18)
I (bases I to 18)
Zwick,M.G., Edington,B.E., McSwiggen,J.A., Merlo,P.Ann.Owens.,Guo,L., Skokut,T.A., Young,S.A., Folkerts,O. and Merlo,D.J.
Nucleic acid encoding delta-9 desaturase
Nucleic us 6350934-A 571 26-PEB-2002;
Location/Qualifiers
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PC A61P35/00,C12N1/15,C12N1/19,C12N1/21,C12NS/10,C12P21/02
PC A61P35/00,C12N9/12,
C12Q1/68/C12N9/12,
PC (A61K35/76,A61K31:522),C12N15/00,A61K37/02,C12N5/00 CC
Description of Artificial Sequence:Oligonucleotide FH Key
Location/Qualifiers
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                               Indels
1. .38
/organism="synthetic construct"
/mol type="genomic DNA"
/db_xref="taxon:32630"
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                                                                                                                                                                               AR196106
Sequence 571 from patent US 6350934.
AR196106.
AR196106.1 GI:20245543
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JP 2002509699-A/40.
                                                               368 AGGAAGAGGAACGGAGC 384
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                                                                                              17 AGGAAAAGGAACGGATC 1
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synthetic construct
              Best Local Similarity 88.2
Matches 15; Conservative
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Best Local Similarity
Matches 15; Conserv
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DEFINITION
ACCESSION
VERSION
KEYWORDS
SOURCE
ORGANISM
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BD225837/c
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JOURNAL
FEATURES
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AUTHORS
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PAT 29-OCT-2001
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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Method and reagent for the inhibition of grid
Parent: WO 0162911-A 857 30-AUG-2001;
RIBOZYME PHARMCEUTICALS, INC. (US); GLAXO GROUP LIMITED (GB)
Location/Qualifiers
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Patent: WO 0188124-A 1481 22-NOV-2001;
RIBOZYME PHARMACEUTICALS, INC. (US) ; GLAXO GROUP LIMITED (GB)
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                                                                                                    Length 17;
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.larity 93.3%; Pred. No. 2.6e+02;
Conservative 0; Mismatches 1;
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Pred. No. 2.6e+02;
0; Mismatches 1;
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Seguence 1481 from Patent WO0188124.
AX423145

    1.17
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/mol_type="unassigned RNA"
/db_xref="taxon:9606"

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/mol_type="unassigned RNA"
/db_xref="taxon:9606"
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Seguence 857 from Patent W00162911.
AX273288
 Location/Qualifiers
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Best Local Similarity 93.3%;
Matches 14; Conservative
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Homo sapiens
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Best Local Similarity
Matches 14; Conserv
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AX423145/c
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Unknown.
Unknown.
Unclassified.
1 (bases 1 to 17)
1 (bases 1 to 17)
Pavco, P., McSwiggen, J.A., Stinchcomb, D.T. and Escobedo, J.
Pavco, P., McSwiggen, J.A., Stinchcomb, D.T. and Escobedo, J.
Patchod and reagent for the treatment of diseases or conditions
related to levels of vascular endothelial growth factor receptor
Patent: US 656127-A 4559 20-MAY-2003;
Location/Qualifiers
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
                                                                                                              Unclassified.

1 (bases 1 to 17)

Guida, M. and Hall, J.

Genetic typing of the human cytochrome P450 2A6 gene and related materials and methods
Patent: US 6492115-A 12 10-DEC-2002;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Jarvis, T., von Carlowitz, I., Mcswiggen, J.A., Hamblin, P.A. and Ellis, J.H.
Method and reagent for the inhibition of grid
Patent: WO 0162911-A 429 30-AUG-2001;
RIBOZYME PHARMACEUTICALS, INC. (US); GLAXO GROUP LIMITED (GB)
                                                                                                                                                                                                                                                                                                                                      Gaps
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Pred. No. 2.6e+02;
0; Mismatches 1; Indels
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US 6566127.
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AX272860
AX272860.1 GI:16545597
Sequence 12 from patent US 6492115.
                                                                                                                                                                                                                                    1. .17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AR327157
Sequence 4559 from patent
AR327157
AR327157.1 GI:33712965
                                                AR264867.1 GI:29693236
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Best Local Similarity 93.3.
Best Local 14; Conservative
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Best Local Similarity 93.3
Matches 14; Conservative
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AX272860/c
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Shay,J.W., Wright,W.E., Piatyszek,M.A., Corey,D. and Norton,J.C.
Modulation of mammalian telomerase by peptide nucleic acids
Patent: JP 2001517929-A 1 09-OCT-2001;
GERON CORP
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                                                           Unclassified.

1 (bases 1 to 13)
Gryaznov, S., Pongracz, K. and Matray, T.
Gryaznov, S., Pongracz, K. and Matray, T.
Oligonuclectide N3', fwdarw, P5' thiophosphoramidates: their synthesis and administration to treat neoplasms
Patent: US 6608036-A 8 19-AUG-2003;
Location/Qualifiers
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Modulation of mammalian telomerase by peptide nucleic acids
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/organism="unidentified"
/mol-type="unassigned RNA"
/db_xref="taxon:32644"
/note="This peptide is synthetically generated"
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2.9%; Score 13; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 2.2e+02;
Matches 13; Conservative 0; Mismatches 0;
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Reduction of hair growth
Patent: WO 03002077-A 3 09-JAN-2003;
THE GILLETTE COMPANY (US)
                                                                                                                                                                                                                                                                                                                                                                                                                                        13 bp
Sequence 3 from Patent WO03002077.
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/organism="unknown"
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BD071035.1 GI:22616638
JP 2001517929-A/1.
unidentified
AR381412.1 GI:40089445
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JP 2001517929-A/1
09-0CT-2001
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Best Local Similarity 100.
Matches 13; Conservative
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BD071035/c
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
                                                                                                                                                                                                                                                                                                      Telerman, A., Amson, R. and Tuijnder, M.
Sequences involved in phenomena of tumour suppression, tumour reversion, apoptosis and/or resistance to viruses and the use thereof as medicaments and 27-MAR-2003;
Molecular Engines Laboratories (FR)
Location/Qualifiers
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1 (bases 1 to 13)

Gryaznov, S., Pongracz, K. and Matray, T.
Oligonucleotide N3'. fwdarw.P5' thiophosphoramidates: their synthesis and administration to treat neoplasms
Patent: US 6608036-A 2 19-AUG-2003;
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                                                                                                                                     linear
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Sequence 1134 from Patent WO03025177.
AX735544
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Sequence 2 from patent US 6608036.
AR381406
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Sequence 8 from patent US 6608036.
AR381412

    .13
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                   124 GGAAAAGCCTCGGCC 138
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                97 TGTTTTTCTCGCTGA 1111
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                                      16 GGAAAAGCCTCCGCC 2
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AX735544/c
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GERON CORP

OS Unidentified

DY 2001517929-A/12

PN UP 2001517929-A/12

PN 09-APR-1997 JP 1997536487

PR 09-APR-1996 US 08/630019

PI JERRY W SHAY, WOODRING E WRIGHT, MIECZYSLAW A PIATYSZEK, DAVID

PI JAMES C NORTON

PC COTKIL4/OO, AGIKS8/16, C12Q1/68

CC Strandedness: Single;

CC Topology: Linear;

CC Topology: Linear;

CC Topology: Linear;

CC Topology: Linear;
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                                                                                                                                                                                                                                                                                                                                                      Shay, J.W., Wright, W.E., Piatyszek, M.A., Corey, D. and Norton, J.C. Modulation of mammalian telomerase by peptide nucleic acids Patent: JP 2001517929-A 12 09-OCT-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Shay, J.W., Wright, W.E., Piatyszek, M.A., Corey, D. and Norton, J.C. Modulation of mammalian telomerase by peptide nucleic acids Patent: JP 2001517929-A 45 09-OCT-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                JAMES C NORTON
C07K14/00,A61K38/16,C12Q1/68
Strandedness: Single;
Topology: Linear;
/desc = 'peptide nucleic acid (PNA), where (deoxy(ribose-phosphate
                                      Gape
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Modulation of mammalian telomerase by peptide nucleic acids.
BD071046
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CC cc nucleotide bases via glycine amino N through a methylenecarbonyl linker'
FH Key location/Qualifiers
FT source longanism='Unidentified'.
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/organism='Unidentified'
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2.9%; Score 13; DB 1; Le
ilarity 100.0%; Pred. No. 2.2e+02;
Conservative 0; Mismatches 0;
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JP 2001517929-A/12.
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JP 2001517929-A/45.
                                                                                                                                                                                                                                                                                  unidentified
unidentified
unclassified.
1 (bases 1 to 13)
                                                                     42 TTGTCTAACCCTA 54
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1 (bases 1 to 13)
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BD071079/c
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Modulation of mammalian telomerase by peptide nucleic acids.
BD071038
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CO7K14/00, A61K38/16, C12Q1/68
Strandedness : Single;
Topology: Linear;
Topology: Linear;
Phosphate
phosphate
                                                                                                             Topology: Linear;
/desc = 'peptide nucleic acid (PNA), where (deoxy(ribose- CC
phosphate
 09-APR-1997 JP 1997536487
09-APR-1996 US 08/630019
JERRY W SHAY,WOODRING E WRIGHT,MIECZYSLAW A PIATYSZEK,DAVID
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Shay.J.W., Wright,W.E., Piatyszek,M.A., Corey,D. and Norton,J.C. Modulation of mammalian telomerase by peptide nucleic acids Pacent: JP 2001517929-A 4 09-OCT-2001;
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FH Key Location/Qualifiers 1. .13
                                                                                                                                                                                                       nucleotide bases via glycine amino N through a lenecarbonyl linker'
                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 2.9%; Score 13; DB 1; Length 13; Best Local Similarity 100.0%; Pred. No. 2.2e+02; Matches 13; Conservative 0; Mismatches 0; Indels
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/organism='Unidentified'
Location/Qualifiers
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/organism='Unidentified'
Location/Qualifiers
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    .13
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/db_xref="taxon:32644"

                                                     COREY,
JAMES C NORTON
CO7K14/00,A61K38/16,C12Q1/68
Strandedness: Single;

    .13
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JP 2001517929-A/4
09-007-2001
09-ARR-1997 JP 1997536487
09-ARR-1996 US 08/630019
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JP 2001517929-A/4.
unidentified
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BD071038/c
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BD071085 13 bp DNA linear PAT 27-AUG-2002 Modulation of mammalian telomerase by peptide nucleic acids. BD071085 BD071085 1 GI:22616688 JP 2001517929-A/51. unidentified
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/desc = 'peptide nucleic acid (PNA), where (deoxy(ribose- CC
phosphate
                                                                                                                                                                                                                                                      1 (bases 1.013)
Shay,J.W., Wright,W.E., Piatyszek,M.A., Corey,D. and Norton,J.C. Modulation of mammalian telomerase by peptide nucleic acids Patent. JP 2001517929-A 51 09-OCT-2001;
                                                                                                                                                                                                                                                                                                                                      Unidentified
JP 2001517929-A/51
09-0CT-2001
09-APR-1997 JP 1997536487
09-APR-1996 US 08/630019
JERRY W SHAY, WOODRING E WRIGHT, MIECZYSLAW A PIATYSZEK, DAVID
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unclassified.
1 (bases 1 to 13)
Shay, J.W., Wright, W.E., Piatyszek, M.A., Corey, D. and Norton, J.C.
Modulation of mammalian telomerase by peptide nucleic acids
Patent: JP 2001517929-A 54 09-OCT-2001;
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CC cc nucleotide bases via glycine amino N through a

methylanecarbonyl linker'

FH Key location/Qualifiers

FT source loganism='Unidentified'.
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Location/Qualifiers
       Pred. No. 2.2e+02;
; Mismatches 0;
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JAMES C NORTON
CO7K14/00,AG1K38/16,C12Q1/68
Strandedness: Single;
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/organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
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      Best Local Similarity 100.0%;
Matches 13; Conservative C
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                                                                  13 TTTTTGTCTAAC 1
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Matches 13; Conservative
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BD071085/c
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BD071084.1 GI:22616687
UP SO0151729-A/50.
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linkages are replaced by N-(2-aminoethyl)glycine units linked
             Unidentified
JP 2001517929-A/45
09-007-2001
09-007-2001
09-APR-1997 JP 1997536487
09-APR-1996 US 08/630019
JERRY W SHAY, WOODRING E WRIGHT, MIECZYSLAW A PIATYSZEK, DAVID
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Unidentified
JP 201517929-A/50
09-CCT-2001
09-APR-1997 JP 1997536487
09-APR-1996 US 08/630019
JERRY W SHAY, WOODRING E WRIGHT, MIECZYSLAW A PIATYSZEK, DAVID
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/desc = 'peptide nucleic acid (PNA), where (deoxy(ribose-CC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   unidentified
unclassified.
1 (bases 1 to 13)
Shay,J.W., Wright,W.E., Piatyszek,M.A., Corey,D. and Norton,J.C.
Modulation of mammalian telomerase by peptide nucleic acids
Patent: JP 2001517929-A 50 09-OCT-2001;
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JAMES C NORTON
COTK14/00, AGIX38/16, C12Q1/68
CO7K14/00, AGIX38/16, C12Q1/68
COTCAGAGES: Single;
C Topology: Linear;
C Topology: Linear;
C Topology: Linear;
Location/Qualifiers
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Strandedness: Single;
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/organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
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JAMES C NORTON
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BD071084/c
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PAT 17-AUG-2003
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(C12NS/00,C12R1:91)
Regulation of repressor genes using nucleic acid molecules FH
Location/Qualifiers
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PN JP 2002541795-A/1700
PP 10-DEC-2002
PF 11-APR-1999 US 60/129390
PP 12-APR-1999 US 60/129390
C12N15/09. AG1K38/00, AG1K48/00, AG1P43/00, C12N5/10, PC C12P21/02,
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Paul, P.S., Meng, X. -J., Halbur, P., Morozov, I. and Lum, M.A.
Polymucleic acids isolated from a porcine reproductive and respiratory syndrome virus (PRRSV) and proteins encoded by the polymucleic acids
Patent: US 6592873-A 34 15-JUL-2003;
Location/Qualifiers
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1 (bases 1 to 17)

1 stt., 2, 2wick, M., Pavco, P. and Mcswiggen, J.

Regulation of repressor genes using nucleic acid molecules Patent: JP 2002541795-A 1700 10-DEC-2002;
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                                                  Query Match 2.9%; Score 13; DB 1; Length 13; Best Local Similarity 100.0%; Pred. No. 2.2e+02; Matches 13; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                 linear
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                                                                                                                                                                                                                                 DNA
                                                                                                                                                                                                                              AR353144 16 bp C
Sequence 34 from patent US 6592873.
AR353144
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /organism="unknown"
/mol_type="genomic DNA"
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                                                                                                                                                                                                                                                                                                                                             Unclassified.
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BD253907/c
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linkages are replaced by N-(2-aminoethyl)glycine units linked
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                                                                                                                                                                  Topology: Linear; /desc = 'peptide nucleic acid (PNA), where (deoxy(ribose- CC
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Unidentified
JP 2001517929-A/54
09-C07-2001
09-APR-1997 JP 1997536487
09-APR-1996 US 08/630019
JERRY W SHAY, WOODRING E WRIGHT, MIECZYSLAW A PIATYSZEK, DAVID
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JP 2001517929-A/55
09-007-2001
09-ARR-1997 JP 1997536487
09-ARR-1996 US 08/630019
JERRY W SHAY,WOODRING E WRIGHT,MIECZYSLAW A PIATYSZEK,DAVID
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Shay.J.W., Wright, W.E., Piatyszek, M.A., Corey, D. and Norton, J.C. Modulation of mammalian telomerase by peptide nucleic acids Patent: JP 2001517929-A 55 09-OCT-2001;
GERON CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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BD071089.1 GI:22616692
JP 2001517929-A/55.
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CC linkages are replaced by N-(2-aminoethyl)glycine
CC cc nucleotide bases via glycine amino N through a methylenecarbonyl linker'
FH Key Location/Qualifiers
FT source location/Qualifiers
FT fource location/Qualifiers
FT fource location/Qualifiers
FT fource location/Qualifiers
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                                                                                                                                                                                                                                                                                                          /organism='Unidentified'

    1. .13
/organism='Unidentified'
Location/Qualifiers

                                                                                              COREY,
JAMES C NORTON
CO7K14/00,A61K38/16,C12Q1/68
Strandedness: Single;

    .13
    /organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"

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JAMES C NORTON
CO7K14/00,A61K38/16,C12Q1/68
Strandedness: Single;
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unclassified.

In (bases 1 to 17)
S Blatt.i., Zantck,M., Pavco,P. and Mcswiggen,J.
Regulation of repressor genes using nucleic acid molecules
Regulation of repressor genes using nucleic acid molecules
RIBOZYME PHARMACHITICALS INC
OS EURARYOLE
PN JP 2002541795-A/6149
PD 10-DEC-2002
PP 11-APR-2000
PP 12-APR-1999 US 60/129390
PR 12-APR-1999 US 60/129390
CI2NIS/09, A6IK38/00, A6IK48/00, A6IP43/00, A6IP43/00, AGIP43/00, AGIP43/00, AGIP43/00, AGIP43/00, AGIP43/00, AGIP43/00, AGIP43/00, AGIPAS/10, PC
                 (bases 1 to 17)
Blasczyk,R.D.
A method for determining the Histocompatibility locus antigen class
                                                                                                                                                                                                                                                                                                                                                                                                                                             BD258356 17-JUL-2003
Regulation of repressor genes using nucleic acid molecules.
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CC Regulation of reserve
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2.8%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 2.8e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
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ilarity 87.5%; Pred. No. 2.8e+02;
Conservative 0; Mismatches 2;
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/organism="unidentified"
/mol_type="unassigned DNA"
/db_xref="taxon:32644"
                                                                                    Patent: EP 0887423-A 3 30-DEC-1998;
BIOTEST AG (DE)
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/organism="unidentified"
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BD258356.1 GI:33068126
JP 2002541795-A/6149.
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BD273168
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BD258356
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DAVID J LOCKHART, THOWAS RYDER, PAMELA SKLAR
C12Q1/68, C12M1/00, C12N15/09, C12N15/09, C12N15/09, G01N33/53, PC
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic construct
synthetic construct
construct
coher sequences; artificial sequences.
1 (bases 1 to 16)
2 Fan, J.B., Hirschhorn, J.N., Huang, X., Kaplan, P., Lander, E.S.,
Lockhart, D.J., Ryder, T. and Sklar, P.
Universal arrays
WHITEHEAD INSTITUTE FOR BIOMEDICAL RESEARCH, AFFYMETRIX INC
OS Artificial Sequence
PN JP 2002539849-A/444
PD 26-NOV-2002
PF 77-MAR-2000 JP 2000608794
PR 26-MAR-1999 US 60/140359 PI
JIAN BING FAN, JOEL N HIRSCHHORN, XIAOHUA
HUANN, PAUL KAPLAN, ERIC
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Location/Qualifiers
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2.8%; Score 12.8; DB 1; Length 16;
Best Local Similarity 87.5%; Pred. No. 2.7e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
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100.0%; Pred. No. 2.7e+02;
tive 0; Mismatches 0; Indels
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PI DAVID J LOCKHART THOMAS RYDER, PAMELA SKLAR
PC C1201/68,C12M1/00,C12N15/09,C12N15/09,C12N15/00
G01N33/566,
PC G01N37/00,C12N15/00,C12N15/00,C12N15/00
CC Primer Location/Qualifiers
FH Key Location/Qualifiers
FT source /organism='Artificial Seque
                         /organism='Eukaryote'.
Location/Qualifiers
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/organism="synthetic construct"
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Sequence 3 from Patent EP0887423.
A82083.
A82083.1 GI:6731948
                                                                            /organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
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JP 2002539849-A/444.
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Best Local Similarity 100.
Matches 13; Conservative
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Universal arrays.
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17 bp
Sequence 27 from Patent W00159103.
AX214585
AX214585.1 GI:15524628
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Sequence 3 from Patent BP0892069.
AX001154
AX001154.1 GI:7241348
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/mol_type="genomic DNA"
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                             GCTGACTTTCAGCGGG
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1 (bases 1 to 17)
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1 (bases 1 to 17)
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Matches 14; Conservative
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AR224417
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AUTHORS
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OS Artificial Sequence
NO 400533113-A/1
PD 08-OCT-2002
PD 08-OCT-2003
PF 23-DEC-1999 JP 2000591168
PR 24-DEC-1999 US 09/220510 PI
NITZAN RESNICK
A61P7/02,
A61P7/02,
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Expression vectors comprising multiple shear stress responsive elements (SSRE) and a gene of interest and methods of use thereof.
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Zwick, M.G., Edington, B.E., McSwiggen, J.A., Merlo, P. Ann. Owens., Guo, L., Skokut, T.A., Young, S.A., Folkerts, O. and Merlo, D.J.
Nucleic acid encoding delta-9 desarurase
Patent: US 6359934-A 828 26-PEB-2002;
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/organism="synthetic construct"
/mol type="genomic DNA"
/db_xref="taxon:32630"
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1 (bases 1 to 17)
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AR196363.1 GI:20245800
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/mol_type="unassigned DNA"
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AR196363
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Expression vectors comprising multiple shear stress responsive elements (SSRE) and methods of use for treating disorders related to vasculogenesis and/or angiogenesis in a shear stress environment Patent: US 6440726-A 1 27-MUG-2002;

Location/Qualifiers
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2.8%; Score 12.8; DB 1;
Best Local Similarity 87.5%; Pred. No. 2.8e+02;
Matches 14; Conservative 0; Mismatches 2;
  DNA
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BIOTEST AG (DE)
AR224417 17 bp
Sequence 1 from patent US 6440726.
AR224417
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Gaps

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PAT 07-SEP-2001

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Fattaey, A.R., Jarvis, T., Mcswiggen, J., Booher, R.N. and Holman, P.S. Method and reagent for the inhibition of checkpoint kinase-1 (chk l) enzyme
Patent: WO 0157206-A 7 09-AUG-2001;
RIBOZYME PHARMACEUTICALS, INC. (US); Fattaey, Ali R. (US)
Location/Qualifiers
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Method and reagent for the modulation and diagnosis of cd20 and nogo gene expression
Patent: WO ALS103-A 1810 16-AUG-2001;
RIBOZYME PHARMACEUTICALS, INC. (US); Blatt, Lawrence (US);
McSwiggen, James (US); Chowrira, Bharat M. (US)
Location/Qualifiers
        nogo gene expression
Patent: WO 0159103-A 59 16-AUG-2001;
RIBOZYME PHARMACEUTICALS, INC. (US); Blatt, Lawrence (US)
McSwiggen, James (US); Chowrira, Bharat M. (US)
Location/Qualifiers
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/ organism="synthetic construct"
/mol_type="unassigned RNA"
/db_xref="texon:32630"
/note="Nucleic Acid"
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/organism="synthetic construct"
/mol_type="unassigned RNA"
/db_xref="taxon:22630"

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    /organism="synthetic construct"
/mol_type="unassigned RNA"

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Sequence 1810 from Patent WO0159103.
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Seguence 7 from Patent WO0157206.
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                                                                                               Blatt, L., Mcswiggen, J. and Chowrira, B.M.
Method and reagent for the modulation and diagnosis of cd20 and nogo gene expression
Patent: WO 0153103-A 27 16-AUG-2001;
RIBOZYME PHARMACEUTICALS, INC. (US); Blatt, Lawrence (US); McSwiggen, James (US); Chowrira, Bharat M. (US)
Location/Qualifiers
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Patent: WO 0159103-A 58 16-AUG-2001,
RIBOZYME PHARMACEUTICALS, INC. (US) ; Blatt,
McSwiggen, James (US) ; Chowrira, Bharat M.
Location/Qualifiers
                                                                                                                                                                                                                                                        /organism="synthetic construct"
/mol type="unassigned RNA"
/db_xref="taxon:32630"
/note="Nucleic Acid"
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/mol_type="unassigned RNA"
/mol_type="unassigned"/db_xref="taxon:32630"
/note="Nucleic Acid"
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                                                              other sequences; artificial sequences.
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Sequence 58 from Patent WO0159103.
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AX214617
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Targeted chromosomal genomic alterations in plants using modified single stranded oligonucleotides
Patent: WO 0195512-A 722 06-DEC-2001;
UNIVERSITY OF DELAWARE (US)
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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Method and reagent for the inhibition of erg
Patent: WO 1018124-A 818 22-NOV-2001;
RIBOZYME PHARMACEUTICALS, INC. (US) ; GLAXO GROUP LIMITED (GB)
Location/Qualifiers
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AX422502.1 GI:21525884
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Sequence 839 from Patent WO0188124.
AX422503
                                                                                17 bp 1
Sequence 752 from Patent W00192512.
AX324614
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CTGAGCTGAGGGCCGT 16
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Best Local Similarity 87.5%;
Matches 14; Conservative
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AX422502/c
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AX324614/c
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Spermatophyta, Magnoliophyta, Liliopsida, Poales, Poaceae, PACCAD
clade, Panicoideae, Andropogoneae, Zea.
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Targeted chromosomal genomic alterations in plants using modified single stranded obligonucleotides
Patent: WO 0195512-A 751 06-DEC-2001;
UNIVERSITY OF DELAWARE (US)
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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Method and reagent for the inhibition of grid
Patent: WO 0162911-A 331 30-AUG-2001;
RIBOZYME PHARMACHICALS, INC. (US) ; GLAXO GROUP LIMITED (GB)
Location/Qualifiers
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87.5%; Pred. No. 2.8e+02;
tive 0; Mismatches 2; Indels
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                                  Query Match
2.8%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 2.8e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
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    /organism="Homo sapiens"
    /mol_type="unassigned RNA"
    /db_xref="taxon:9606"

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/db_xref="taxon:4577"
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/db xref="taxon:32630"
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/organism="Zea mays"
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AX272762/c
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AX324613
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PAT 26-NOV-2002

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Homo sapiens
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Human udp-galnac:polypeptide n-acetylgalatosaminyltransferase 10
Patent: EP 1243660-A 739 25-SEP-2002;
Aeomica, Inc. (US)
Location/Qualifiers
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Abang,J., Gu,Y. and Nguyen,C.T.
Human udp-galnac:polypeptide n-acetylgalatosaminyltransferase 10
Patent: EP 1243660-A 738 25-SEP-2002;
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87.5%; Pred. No. 2.8e+02;
tive 0; Mismatches 2; Indels
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Best Local Similarity 87.5%; Pred. No. 2.8e+02;
Matches 14; Conservative 0; Mismatches 2;
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Sequence 739 from Patent EP1243660.
AX545226

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/mol_type="unassigned DNA"
/db_xref="taxon:9606"

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Sequence 738 from Patent EP1243660.
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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Homo sapiens
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Best Local Similarity 87.55
Matches 14; Conservative
Inc. (US)
   Aeomica,
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AX545226/c
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AXS45225/c
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                           Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi,
Mammalia, Butheria, Primates, Catarrhini, Hominidae, Homo.
                                                                                                                         Jarvis,T., von Carlowitz,I., Mcswiggen,J.A., Mclaughlin,F.G. a Randi,A.M.
Method and reagent for the inhibition of erg
Patent: WO 0188124-A 839 22-NOV-2001;
RIBOZYME PHARMACEUTICALS, INC. (US) ; GLAXO GROUP LIMITED (GB)
Location/Qualifiers
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Patent: EP 1239051-A 716 11-SEP-2002;
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Patent: BP 1239051-A 715 11-SEP-2002;
Aeomica, Inc. (US)
Location/Qualifiers
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Sequence 716 from Patent EP1239051.
AX531207.
AX531207.1 GI:25254207
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       AX422503.1 GI:21525885
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Kit and method for determining HLA type
Patent: WO 019257-2A 1071 06-DEC-2001;
NISSHINBO INDUSTRIES INC, SYSTEM RESEARCH INC, HIDETOSHI INOKO, TAEKO
KAGIYA, TATSUO ICHIHARA,YOSHIYUKI MATSUMURA,SHOGO MORIYA,MICHIO
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Patent: JP 2002509699-A 44 02-APR-2002;
CANCER RESEARCH CAMPAIGN TECHNOLOGY LTD
OS Artificial Sequence
PN JP 2002509699-A/44
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                                        synthetic construct
synthetic construct
other sequences; artificial sequences.
1 (bases 1 to 17)
Inoko,H., Kagiya,T., Ichihara,T., Matsumura,Y., Moriya,S. and
Mishida,M.
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01-JUN-2001 WO 2001JP004662
01-JUN-2000 JP 00P 164798
HIDETOSHI INOKO, TAEKO KAGIYA, TATSUO ICHIHARA, YOSHIYUKI
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Location/Qualifiers
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C12Q1/68,C12M1/00,C12N15/09,G01N33/53
Description of Artificial Sequence:capture
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1 (bases 1 to 25)
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29-JAN-1998 GB 9801902.9
WILLIAM NICOL KEITH
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WO 0192572-A/1071
BD104967.
BD104967.1 GI:22650541
WO 0192572-A/1071.
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PC A61P35
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PF 01-JUN
PI HIDETC
MATSUMURA,
PI SHOGO
PC C1201/
CC Descri
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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Four human zinc-finger-containing proteins : mdz3, mdz4, mdz7 and
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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17 bp DNA
Kit and method for determining HLA type.
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Aeomica, Inc. (US)
Location/Qualifiers
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Aeomica, Inc. (US)
Location/Qualifiers
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Sequence 815 from Patent EP1281758.
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Sequence 813 from Patent EP1281758.

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   343 GCGAGGTTCAGGCCTT 358
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                                   16 GCGCGGATCAGGCCTT
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BD023698 30 bp DNA linear PAT 27-AUG-2002 Method for detecting and inhibiting RNA component of telomerase.
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Promoter region of mouse and human telomerase RNA component genes.
BD225839
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Kim,N.W., Wu,F., Kealey,J.T., Pruzan,R. and Weinrich,S.L.

Method for detecting and inhibiting RNA component of telomerase

Patent: JP 2001507229-A 2 05-JUN-2001;

GERON CORP
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PC A61P35/00,C12N1/15,C12N1/19,C12N1/21,C12N5/10,C12P21/02 PC
C1201/68/C12N9/12,
PC (A61K315/76,A61K31:S22),C12N15/00,A61K37/02,C12N5/00 CC
Description of Artificial Sequence:Oligonucleotide FH Key
                                                                                                                                                                                                                                                                                                     PN JP 2001507229-A/2
PD 05-UTN-2001
PF 19-DEC-1997 JP 1998529003
PR 20-DEC-1996 US 08/770554,20-DEC-1996 US 08/770565 PI
NAM WOO KIM, FRED WU, JAMES T KEALEY, RONALD PRUZAN, SCOTT L PI
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Pred. No. 3.6e+02;
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synthetic construct
other sequences; artificial sequences.
1 (bases 1 to 38)
Keith, W.N.
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Best Local Similarity 70.8%; Pred. No. 3.6e
Matches 17; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
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                                                                                                                       BD023698.1 GI:22564921
JP 2001507229-A/2.
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unclassified
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VERSION
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VERSION
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Km,N,N,Oo., Wu,F., Kealey,J.T., Pruzan,R. and Weinrich,S.L.
Methods for detecting the RNA component of telomerase
Patent: US 5846723-A 2 08-DEC-1998;
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KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH (GB)
Location/Qualifiers
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                                                                                                        Length 25;
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    .25
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|db_xref="taxon:32630"

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/organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="teaxon:32630"
/noTe="Oligonucleotide"
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other sequences; artificial sequences.
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/organism="unknown"
/mol_type="unassigned DNA"
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Sequence 2 from patent US 5846723.
AR063826.
AR063826.1 GI:5993134
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Sequence 44 from Patent WO9938964.
                                                                                                                                                                               223 CGCCTGCCCAGCCCCCGAACCCCG 246
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AX019590.1 GI:10043504
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Best Local Similarity 70.8%;
Matches 17; Conservative
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AX019590/c
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AR063826
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Roninson, I.B. and Grossman, A.
Compositions, methods and kits for identifying naturally occurring
RNA sequences having affinity for RNA-binding proteins
Patent: US 6238867-A 15 29-MAY-2001;
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PR 29-JNN-1998 GB 9801902.9
PI WILLIAM NICOL KEITH
PC CI2N15/09, A61K31/7105, A61K31/711, A61K35/76, A61K38/00, A61K45/00, PC
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UP 200250940.0 AI:33035650
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Promoter region of mouse and human telomerase RNA component genes
Patent: JP 200259659-A 83 02-APR-2002;
CANCER RESEARCH CAMPAIGN TECHNOLOGY LTD
OS Artificial Sequence
PN JP 2002509699-A/83
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PC A61P35/00,C12N1/15,C12N1/19,C12N1/21,C12N5/10,C12P21/02 PC
.C12Q1/68//C12N9/12.
PC (A61K35/76,A61K31:522),C12N15/00,A61K37/02,C12N5/00 CC
Description of Artificial Sequence: Mutant construct FH Key
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                                                                                                                               linear
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synthetic construct
synthetic construct
other sequences; artificial sequences.
I (bases 1 to 51)
                                                                                                                             AR154150 66 bp I
Sequence 15 from patent US 6238867.
AR154150.
GI:15122203
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29-JAN-1999 JP 2000529424
29-JAN-1998 GB 9801902.9
WILLIAM NICOL KEITH
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Unclassified.
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BD225880/c
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Roninson, I.B. and Grossman, A.
Compositions, methods and kits for identifying naturally occurring
RNA sequences hathod affinity for RNA-binding proteins
Patent: US 6238867-A 6 29-MAY-2001;
Location/Qualifiers
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KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH (GB)
Location/Qualifiers
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                                                                                                                                               Query Match 2.8%; Score 12.8; DB 1; Length 38; Best Local Similarity 70.8%; Pred. No. 3.3e+02; Matches 17; Conservative 0; Mismatches 7; Indels
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/noTe="Oligonucleotide"
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/mol_type="genomic DNA"
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Sequence 16 from patent US 6238867.
AR154151 dr:15122204

    . 62
    /organism="unknown"
    /mol_type="unassigned DNA"

                                                                                                                                                                                                                                                                                                                               Sequence 42 from Patent W09938964. AX019588
                                                                                                                                                                                                                   223 CGCCTGCCCAGCCCCGAACCCCG 246
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Best Local Similarity 70.8
Matches 17; Conservative
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Best Local Similarity
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Promoter region of mouse and human telomerase RNA component genes.
BD225882
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C12Q1/68//C12N9/12,

PC (AGIN35/76,AGIN31:522),C12N15/00,AG1N37/02,C12N5/00

Description of Artificial Sequence: Mutant construct FH

Location/Qualifiers
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KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH (GB)
Location/Qualifiers
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60.0%; Pred. No. 2.7e+02;
tive 0; Mismatches 14;
                                            263 GGCCCGGGGCTTCTCCGGAGGCACCCACTGCCACC 297
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60.0%; Pred. No. 2.7e+02;
tive 0; Mismatches 14;
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                                                                                                                                                 Sequence 83 from Patent W09938964.
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29-JAN-1998 GB 9801902.9
WILLIAM NICOL KEITH
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Best Local Similarity 60.0%
Matches 21; Conservative
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AX019629/c
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Sullivan, S., Draper, K., Kisich, K., Stinchcomb, D.T. and McSwiggen, J. TNF-.alpha. ribozymes
Patent: US 5811300-A 658 22-SEP-1998;
Location/Qualifiers
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KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH (GB)
Location/Qualifiers
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2.8%; Score 12.6; DB 1; Length 51;
Best Local Similarity 60.0%; Pred. No. 2.7e+02;
Matches 21; Conservative 0; Mismatches 14; Indels
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|mol_type="unassigned DNA"
|bxref="taxon:32630"
|noE="Mutant construct"
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AR041868
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Sequence 85 from Patent W09938964.
AX019631
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/wol_type="unassigned DNA"
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Location/Qualifiers
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Best Local Similarity 92.9
Matches 13; Conservative
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L Submitted (21-SEP-2004) Balzergue S., UMRGV, INRA/CNRS, 2 rue

Gaston Cremieux, 91057 Evry cedex, FRANCE
Gaston Cremieux, 91057 Evry cedex, FRANCE

PCR was performed on DNA from transformants of Arabidopsis thaliana
plants from INRA (Versailles). The DNA fragment (s) resulting from
the PCR were directly sequenced from the left or the right border
to determine the genomic sequence flanking the insertion. T-DNA
derived sequences were removed. Information to order the
corresponding mutant line and a link to a database providing a
graphical display of the insertion site are available at
http://dbsgap.versailles.inra.fr/publiclines/. This sequence has
been generated in the framework of the French plant genomics
program 'Genoplante' (http://www.genoplante.com and
http://genoplante-info.infobiogen.fr).
                                        PAT 21-FEB-2003
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T-DNA integration into the Arabidopsis genome depends on sequences
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rosids, eurosids II, Brassicales, Brassicaceae, Arabidopsis.
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Arabidopsis thaliana (thale cress)
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                              Sequence 4425 from Patent EP1260586. AX637286 GI:28472900

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/clone_lib="Arabidopsis thaliana T-DNA insertion lines"
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Method of diagnosing colon and gastric cancers
Patent: WO 2004021010-A 81 11-MAR-2004;
Oncotherapy Science, Inc. (JP); Japan as represented by the president of the university of Tokyo (JP)
Location/Qualifiers
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| Mol_Lype="unassigned DNA"
| db xrefe="taxon.32630"
| /noTe="Artificially synthesized S-oligonucleotide"
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Guida,M. and Kurth,J.
Genotyping the human phenol sulfotransferbase 2 gene STP2
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                 /mol_type="genomic DNA"
/cultivar="Wassillewskija"
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Keith, T.

Novel human gene relating to respiratory diseases, obesity, and inflammatory bowel disease

L Patent: WO 017884-A 316 25-OCT-2001;

Genome Therapeutics Corp. (US)

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/ organism="synthetic construct"
//mol type="unassigned DNA"
//db xref="taxon:32630"
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Hepatitis B virus 3671 HCV DNAzyme substr HCR COLL HCR CALL H	T 1 647 ADP27647 standard; DNA; 79 BP. ADP27647 standard; DNA; 79 BP. 26-AUG-2004 (first entry) Human TERC DNA used as a cancer prognostic marker SeqID 84. TERC; human; PCR amplicon; ds; prognostic marker; EGPR; epidermal growth factor receptor; cancer; gene expression profiling; microarray; head and neck cancer; colon cancer; metastatic spread; homo sapiens. WO2004046386-A1.	WOZDOW4046386-A1. 13-JUN-2004. 14-NOV-2003; 2003WO-US036777. 15-NOV-2002; 2002US-042709DP. (GENO-) GENOMIC HEALTH INC. (VALL-) VALL HEBRON UNIV HOSPITAL. Baker JB, Cronin MT, Shak S, Baselga J; Prognosing a patient with EGFR-expressing colon cancer comprises subjecting a sample comprising EGFR-expressing cancer cells to quantitative analysis of the expression level of the RNA transcript of least one gene e.g., CD44v3. This invention relates to a novel method concerning prognostic markers associated with EGFR (epidermal growth factor receptor) positive cancer. Specifically, it refers to a gene expression profiling method that can provide a prediction as to agene expression profiling method that can provide a prediction as to whether a patient is likely to respond well to treatment with an EGFR inhibitor. The present invention describes the quantitative analysis of the expression level of the RNA transcript of the est product the group of CD44v3, CD44v6, RR5, GROI, KRT17, LAMC2 or their products thereof. It further provides a CDNA which are useful for determining whether a patient diagnosed with an EGFR expressing head or neck cancer or colon cancer exhibits elevated or beckpressing head or neck cancer or colon cancer exhibits elevated or
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6 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8	AL. 1647 ADP27647 standard; DNA; 79 BP. ADP27647; 26-AUG-2004 (first entry) Human TERC DNA used as a cancer pi TERC; human; PCR amplicon; ds; pri microarray; head and neck cancer; neoplastic disease. Homo sapiens.	WOZUGU4046386-A1. 03-JUN-2004. 14-NOV-2003; 2003WO-US036777. 15-NOV-2002; 2002US-0427090P. (GENO-) GENOMIC HEALTH INC. (VALL-) VALL HEBRON UNIV HOSPITAL. Baker JB, Cronin MT, Shak S, Baselga Prognosing a patient with EGFR-expression expression of the expression of the expression least one gene e.g., CD44V3. Claim 55; SEQ ID NO 84; 113pp; English. This invention relates to a novel methor associated with EGFR (epidermal growth Specifically, it refers to a gene expression expression with an EGFR inhibitor. The provide a prediction as to whether a preatment with an EGFR inhibitor. The provide a prediction as to whether a prediction as to whether a pression energene selected from the group KRTI7, LAMC2 or their products thereof. Which are useful for determining whether expressing head or neck cancer or colodecreased expression levels of these egenerates.
C C C C C C C C C C C C C C C C C C C	SUL.	PN WOZOG PD 75 14-NO XX 15-NO XX 15-NO PR 15-NO PR (GENO PR (WPL; XX XX YALLA XX YALLA XX Claim XX Claim
Match 2, Telomera Telomera Telomera Labelled Labelled Human G, 2,5,-01 Calanin Human LC Ruman CG Human LC Ruman CG Ruman HO Human HO Human HO Human HO Oligo HC Reverse Inhibito	Seroconin Partiviral Antiviral Delta-9 de Human BRCA Group-spec Human leuk (/BBP-bete Human aden Human (ARI Human Chki Human (OKI Human (OKI Human (OKI Human NOGC Human NOGC	MALE-SEC MALE-SEC MALE-SEC HUMAN F HUMAN F HUMAN F HUMAN H WAV AIN WAV AIN WAV AIN WAV AIN WAV HUMAN H HUMAN I H H HUMAN I H H H H H H H H H H H H H H H H H H H
ADB68046 ABZS8497 ADM46660 AD021607 AD021507 AD021807 AD021807 AAV89815 AAV89814 ABK97507 AAK97507	24	1999 1997 1997 1997 1997 1997 1997 1997
	ADN13 AAT1 AAT1 AAX0 AAX0 AAX0 AAX0 AAX0 AAX1 AAA7 AAA7 AAA7 AAH9 AAH9 ABK0 ABK0 ABK0 ABK0 ABK0 ABK0 ABK0 ABK0	ABK25391 ABK18192 ABK18192 ABK18191 ABV90002 ABU31582 ACN04592 ACN04059
		17 1 ABK25 17 1 ABK25 17 1 ABK18 17 1 ABV90 17 1 ABV90 17 1 ACN04 17 1 ACN04 17 1 ACN03 17 1 ACN03 17 1 ABZ65 17 1 ABZ61 17 1 ABZ61
133 2.9 133 133 2.9 133 2.9 133 2.9 133 2.9 133 2.9 133 2.9 133 2.9 134 133 2.9 144 133 2.9 155 133 2.9 155 133 2.9 155 133 2.9 155 133 2.9 155 134 2.8 156 2.	8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8	22.22.22.23.88.88.88.88.88.88.88.88.88.88.88.88.88

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invention

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detecting a first subunit protein (1) of telomerase. R1 and R2 both bind to (1) and have formulae 5'-A-B-C-3' (R1) 5'-D-E-F-3' (R2) where: A and F c R not and a segment of 10 to 100000 nucleotides (nt) that together are replicated by RNA replicase; B and E = RNA segments of 10 to 250 nt from the Y region of human telomerase and bind specifically to (1); C and D = RNA segments of 1 to 10000 nt which can be ligated together. Ligation of RNA segments of 1 to 10000 nt which can be ligated together. Ligation of to (1). Replication of R3 by RNA replicase indicates presence of (1). Also described are: (1) method for detecting (I) using R1 and R2; (2) kit for this process containing R1, R2, by transcription from appropriate DNA. R1 and R2 are used to detect and quantify telomerase, particularly cordiagnosis of cancer and for detection of metastases. R1 and R2 present assay that does not require expensive equipment or highly trained personnel, and is suitable for automation. The present sequence represent an oligonucleotide used in the exemplification of the present
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these methods are also useful for prognosing or predicting the likelihood of cancer-attributable death or progression, including recurrence and metastatic spread of a neoplastic disease, as well as drug resistance. This polynuclectide sequence is a human PCR amplicon DNA sequence used as a prognostic cancer marker, given in an exemplification of the invention.
                                                                                                                                                                                                                                  371 AAGAGGAACGGAGCGAGTCCCCGCGCGCGCGCGATTCCCTGAGCTGTGGGACGTGCACC 430
                                                                                                                                                                                                                                                        1 AAGAGGAACGGAGCGAGCGCGCGCGCGCGATTCCCTGAGCTGGGGACGTGCACC 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pair of RNA molecules for detecting telomerase, useful for diagnosis of cancer or metastases, can be ligated when bound to telomerase subunit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human, telomerase, hTR, telomeric repeat amplification protocol; TRAP,
identification, detection; quantification, cancer, metastasis, ss.
                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention describes a pair of RNA molecules (R1, R2) for
                                                                                                                                                                                              ö
                                                                                                                                                        Length 79;
                                                                                                                                                                                            0; Indels
                                                                                                                  Sequence 79 BP; 16 A; 26 C; 27 G; 10 T; 0 U; 0 Other;
                                                                                                                                                      Score 79; DB 1; I
Pred. No. 0.0059;
                                                                                                                                            17.5%; Sco. 100.0%; Pred. No. v. 100.0%; Pred. No. v. 100.0%; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 2; Col 23; 32pp; English.
                                                                                                                                                                                                                                                                                                          431 CAGGACTCGGCTCACACAT 449
                                                                                                                                                                                                                                                                                                                                  CAGGACTCGGCTCACACAT 79
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                                                                                                                                                                                                                                                                                                                                                                                                                                               AAA08205 standard; DNA; 62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                          79; Conservative
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                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mastadenovirus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12-JUN-1997;
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components. RNA1 has the formula 5'-A-B-C-3', where A is section having components. RNA1 has the formula 5'-A-B-C-3', where A is section having components. RNA1 has the formula 5'-A-B-C-3', where A is section having components. RNA1 has the formula 5'-A-B-C-3', where A is section having components. RNA1 has the formula 5'-A-B-C-3', where A is section having 10-3,000 nucleotides which have affinity to one consensus sequence of the RBP and which can bind to the protein component; C is section having a about 1-20 nucleotides which have affinity to second consensus sequence of the RBP conclectides which have affinity to second consensus sequence of the RBP conclectides which have affinity to second consensus sequence of the RBP conclectides which have affinity to second consensus sequence of the RBP conclectides which have affinity to second consensus sequence of the RBP conclectides which have affinity to second consensus as sequence of the RBP conclectides which can be received by an RNA replicase and with A being replicated. RNA1 and RNA2 are capable of forming a third RNA (RNA3) coff formula 5'-B-E-F-3'. The method is useful for the identification and characterization of RNA sequences having specific affinity to amino acid consensus sequences of RBPs, and to RNAs. AAH24815-16 were used to be a consensus acused to RBPs, and to RNAs. AAH24815-16 were used to be a consensus sequence a double-stranded RNA1, comprising the Y-1 domain of human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New ribonucleic acids useful for identifying naturally occurring RNA sequences having affinity for RNA-binding protein having protein and RNA
                                                                                                                                                                                                                                                                                                                                                                                                          RNA-binding protein; RBP; RNA replicase; RNA identification; telomerase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The specification describes a first RNA (RNA1) and a second RNA (RNA2)
                                                                            Gaps
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                                                                                                                 54
                                                                                                                                                     8
                                                                                                                                                                                                                                                                                                                                                                     Human nucleic acid sequence derived from Y-1 domain of telomerase.
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                                        DB 1; Length 62;
0.7;
                                                                            IndelB
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Sequence 62 BP; 16 A; 27 C; 10 G; 9 T; 0 U; 0 Other;
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100.0%; Pred. No. v...
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Conservative 0; Mismatches
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                                                                                                                                                                                                                                                    AAH24816 standard; RNA; 62
                                                                                                                                                                                                                                                                                                                                 (first entry)
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                                                                              54; Conservative
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es' 54; Conserv
                                        Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US6238867-B1.
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                                                                                                                                                                                                                                                                                           AAH24816;
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                                                                                                                                                                                                                RESULT 3
AAH24816/c
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Matches'
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Matches
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RNA-binding protein; RBP; RNA replicase; RNA identification; telomerase;

Homo sapiens. JS6238867-B1.

Human nucleic acid sequence derived from Y-1 domain of telomerase.

AAH24815 standard; RNA; 66 BP.

06-AUG-2001

AAH24815;

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The specification describes a first RNA (RNA1) and a second RNA (RNA2) for use in binding an RNA-binding protein (RBP) having protein and RNA components. RNA1 has the formula 5'.A-B-C-3', where A is section having (10-100,000 nucleotides and is can be received by an RNA replicase and with another DNA sequence, F, being replicated; B is section having 10-3,000 nucleotides which have affinity to one consensus sequence of the RBP and which can bind to the protein component; C is section having about 1-20 nucleotides and which can be ligated to D of the second RNA molecule. RNA2 has the formula 5'.D-E-F-3', where D is section having 1-C and uncleotides and which can be ligated to C; E is section having 1-C and which can bind to the protein component; F is section having 10-100,000 nucleotides which have affinity to second consensus sequence of the RBP and which can bind to the protein component; F is section having 10-100,000 nucleotides which can be received by an RNA replicase and with A lobeing replicated. RNA1 and RNA2 are capable of forming a third RNA (RNA3) of formula 5'A-B-C-D-E-F-3'. The method is useful for the identification of A characterization of RNA sequences having specific affinity to anino
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention describes a pair of RNA molecules (R1, R2) for detecting a first subunit protein (1) of telomerase. R1 and R2 both bind to (1) and have formulae 5'-A-B-C-3' (R1) 5'-D-E-F-3' (R2) where: A and F - RNA segment of 10 to 100000 nucleotides (nt) that together are replicated by RNA replicase; B and E = RNA segments of 10 to 250 nt from the Y region of human telomerase and bind specifically to (1); C and D = RNA segments of 1 to 10000 nt which can be ligated together. Ligation of C and D produces R3 of formula 5'-A-B-C-D-E-F-3' (R3) with E and B bound to (1). Replication of R3 by RNA replicase indicates presence of (1). Also described are: (1) method for detecting (1) using R1 and R2; (2) kit for this process containing R1, R2, ligase and an amplification system; and (3) method for making R1 and R2 by transcription from appropriate DNA. R1 and R2 are used to detect and quantify telomerase, particularly for diagnosis of cancer and for detection of metastases. R1 and R2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  provide an assay that does not require expensive equipment or highly trained personnel, and is suitable for automation. The present sequence represents an oligonucleotide used in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pair of RNA molecules for detecting telomerase, useful for diagnosis of cancer or metastases, can be ligated when bound to telomerase subunit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; telomerase; hTR; telomeric repeat amplification protocol; TRAP; identification; detection; quantification; cancer; metastasis; ss.
                            54
                                                                              GGGTTGCGGAGGGTGGGCCTGGGAGGGTGGTGGCCATTTTTTGTCTAACCCTA 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12.0%; Score 54; DB 1; Length 66; 100.0%; Pred. No. 0.77;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 66 BP; 10 A; 11 C; 28 G; 17 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                        Adenovirus nucleotide sequence SEQ ID NO:19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 2; Col 23; 32pp; English.
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                                                                                                                                                                                                                                        AAA08204 standard; DNA; 66 BP
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                                                                                                                                                                                                                                                                                                 AAA08204;
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New ribonucleic acids useful for identifying naturally occurring RNA sequences having affinity for RNA-binding protein having protein and RNA

(INVI-) INVITRO DIAGNOSTICS INC.

WPI; 2001-366472/38.

Roninson IB,

99US-00255464 98US-0075495P

22-FEB-1999; 23-FEB-1998;

29-MAY-2001.

Example 2; Col 26; 36pp; English.

components.

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acid consensus sequences of RBPs, and to RNAs. AAH24815-16 were used to produce a double-stranded RNA1, comprising the Y-1 domain of human
                                                                                                                                                           Gaps
                                                                                                                                                                                                1 GGGTTGCGGAGGGTGGCCTGGGAGGGGTGGTGGCCATTTTTGTCTAACCCTA
                                                                                                                                                                                                                                  GGGTTGCGGAGGGTGGAGGGGTGGTGGCCATTTTTGTCTAACCCTA
                                                                                                                                                                                                                                                                                                                                                                                                                                          Mutated hTR promoter fragment containing construct 29111 (mRCE).
                                                                                                                                                           ;
0
                                                                                                                     12.0%; Score 54; DB 1; Length 66; 100.0%; Pred. No. 0.77;
                                                                                                                                                           0; Indels
                                                                                Sequence 66 BP; 10 A; 11 C; 28 G; 17 T; 0 U; 0 Other;
                                                                                                                             100.0%; Prea. ....
                                                                                                                                                                                                                                                                                                                                BP.
                                                                                                                                                                                                                                                                                                                              AAZ00334 standard; DNA; 51
                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                           54; Conservative
                                                                                                                                       Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                       22-OCT-1999
                                                                                                                                                                                                                                                                                                                                                                   AAZ00334;
                                                                                                                       Query Match
                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                         RESULT 6
AAZ00334
ID AAZ0
XX
AC AAZ0
XX
DT 22-O
XX
DE Muta
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.; 0

Gaps

0; Indels

100.0%; Pred no.

54; Conservative

Matches

6 q RESULT 5 AAH24815

Local Similarity

59

1 GGGTTGCGGAGGGTGGGCCTGGGAGGGGTGGTGGCCATTTTTTTGTCTAACCCTA

RES CAMPAIGN TECHNOLOGY.

99WO-GB000308 98GB-00001902

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Mouse and human telomerase RNA gene promoters, useful for tumor specific
                                                                                                                                                                  WPI; 1999-479183/40.
                                                                                                                (CANC-) CANCER
             409938964-A2
                                                               29-JAN-1999;
                                                                                                                                                                                                           gene therapy
                                     05-AUG-1999
                                                                                                                                         Keith WN;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 8
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                                                                                                                                                                                                                                                                                                                                 The invention relates to promoter regions from mouse and human telomerase

RNA (TR) component genes. The TR gene promoter can be linked to a

heterologous gene, especially a gene encoding a cytotoxin, for therapy of

cancer, especially neoplasias. The telomerase is necessary for the

currestricted proliferative capacity of many human cancers. Mutation or

curregulation of the telomerase repression pathway may cause reactivation

or upregulation of telomerase expression in cancer. Substances,

clene promoter through interaction of the 5' regulatory sequences. These

substances, e.g. antisense oligonucleotides, transcription factors,

cuseful for cancer therapy. In particular, gene therapy vectors

cuseful for cancer therapy. In particular, gene therapy vectors

(sepecially pGT62-codAupp) comprising the promoter and a viral thymidine

kinase gene can be used to convert a prodrug, e.g. gancyclovir, so that

ceplasia can be controlled or treated. Direct down-regulation of

telomerase RNA gene through manipulation of transcription factors may be

effective anticancer therapy and the cloning of the hTR gene promoter

allows the analysis of therapeutic molecules which modulate hTR promoter

cativity. Sequences AAZ07324-332 and AAZ0032-340 represent constructs

with hTR promoter sequence element mutations
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                                                                                                                                                                                                                                                                        and human telomerase RNA gene promoters, useful for tumor specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
            Telomerase RNA; TR; promoter; cytotoxin; cancer; neoplasia; hTR;
gene therapy; thymidine kinase gene; anticancer therapy; human; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Telomerase RNA; TR; promoter; cytotoxin; cancer; neoplasia; hTR; gene therapy; thymidine kinase gene; anticancer therapy; human; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mutated hTR promoter fragment containing construct 29112 (mSpl.4).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 GGGTTGCGGAGGGTGGCCTGGGAGGGGTGGTGGCCATTTTTTGTCTAACC 51
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GGTTGCGGAAAATGGGCCTGGGAGGGGTGGTGGCCATTTTTGTCTAACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10.2%; Score 46.2; DB 1; Length 51; 94.1%; Pred. No. 2.6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 51 BP; 8 A; 8 C; 21 G; 14 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
                                                                                                                                                                                             (CANC-) CANCER RES CAMPAIGN TECHNOLOGY.
                                                                                                                                                                                                                                                                                                              Disclosure; Fig 19; 109pp; English
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                                                                                                                                         99WO-GB000308
                                                                                                                                                                   98GB-00001902
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les 48; Conservative
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                                                                                                                                                                                                                                               WPI; 1999-479183/40.
                                                                                                                                                                                                                                                                                       gene therapy.
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Synthetic.
                                                 sapiens
                                                                                      WO9938964-A2
                                                                                                                                         29-JAN-1999;
                                                                                                                                                                   29-JAN-1998;
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                                                                                                                05-AUG-1999.
                                                               Synthetic.
                                                                                                                                                                                                                      Keith WN;
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                                                                                                                                                                                                                                                                          Mouse
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Matches
                                                    Homo
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The invention relates to promoter regions from mouse and human telomerase
RNA (TR) component genes. The TR gene promoter can be linked to a
Cheterologous gene, especially a gene encoding a cytotoxin, for therapy of
Cancer, especially neoplasias. The telomerase is necessary for the
Unrestricted proliferative capacity of many human cancers. Mutation or
dysregulation of the telomerase expression pathway may cause reactivation
Cor upregulation of telomerase expression pathway may cause reactivation
Cor upregulation of telomerase expression pathway may cause reactivation
Cor upregulation of telomerase expression pathway may cause reactivation
Cor upregulation of telomerase expression pathway and reactivation
Cor upregulation of telomerase expression in cancer. Substances,
cleaped acceptances, e.g. antisense oligonucleotides, transcription from the TR
Corpecially proffed-coddupp) comprising the promoter and a viral thymidine
Conspecially proffed-coddupp) comprising the promoter and a viral thymidine
Corplasia can be used to convert a product down regulation of
Confective anticancer therapy and the cloning of transcription factors may be
Coffective anticancer therapy and the cloning of the hTR gene promoter
Coffective anticancer therapy and the cloning of the hTR gene promoter
Coffective anticancer therapy and the cloning of the hTR gene promoter
Coffective anticancer therapy and the cloning of the hTR gene promoter
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            51
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            with hTR promoter sequence element mutations
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Pred. No. 2.6;
0; Mismatches
Disclosure; Fig 19; 109pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAZ00337 standard; DNA; 47 BP
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Best Local Similarity
Matches 48; Conserv
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Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAZ00339
δ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                원
                                                                                                                                                                                   RMA (TR) component genes. The TR gene promoter can be linked to a heterologous gene, especially a gene encoding a cytotoxin, for therapy of cancer, especially neoplasias. The telomerase is necessary for the unrestricted proliferative capacity of many human cancers. Mutation or dysregulation of the telomerase repression pathway may cause reactivation or dysregulation of the telomerase expression pathway may cause reactivation or dentified in the methods, can be used to block transcription from the TR gene promoter through interaction of the 5' regulatory sequences. These substances, e.g. antisense oligonucleotides, transcription factors, peptide nucleic acids and factors that disrupt signal transduction, are useful for cancer therapy. In particular, gene therapy vectors (especially pGT62-codAupp) comprising the promoter and a viral thymidine kinase gene can be used to convert a prodrug, e.g. gancyclovir, so that neoplasia can be controlled or treated Direct down-regulation of telomerase RNA gene through manipulation of transcription factors may be effective anticancer therapy and the cloning of the hTR gene promoter allows the analysis of therapeutic molecules which modulate hTR promoter activity. Sequences AAZ07324-332 and AAZ00332-340 represent constructs
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                                                                                                                                                                          The invention relates to promoter regions from mouse and human telomerase
                                                                                                             Mouse and human telomerase RNA gene promoters, useful for tumor specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Telomerase RNA; TR; promoter; cytotoxin; cancer; neoplasia; hTR; gene therapy; thymidine kinase gene; anticancer therapy; human; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mutated hTR promoter fragment containing construct 111(mSpl.4).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1; Length 47;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GTGGAGGGTGGGTAAGGTGGTGGCCATTTTTTGTCTAACC 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GCGGAGGGTGGCCTGGGAGGGGTGGTGGCCATTTTTGTCTAACC
                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 47 BP; 6 A; 7 C; 20 G; 14 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                    with hTR promoter sequence element mutations
                                                                                                                                                                                                                                                                                                                                                                                                                                                                8.8%; Score 39.6; DE
31.3%; Pred. No. 8.9;
ive 0; Mismatches
                                    (CANC-) CANCER RES CAMPAIGN TECHNOLOGY.
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                                                                                                                                                  Disclosure; Fig 19; 109pp; English
             98GB-00001902.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity 91.3%;
les 42; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAZ00336 standard; DNA; 47
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                                                                                     WPI; 1999-479183/40
             29-JAN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29-JAN-1998;
                                                                                                                            gene therapy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAZ00336;
                                                             Keith WN;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 9
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The intention features to promoter regions from a consistences.

And (TR) component genes. The TR gene encoding a cytotoxin, for therapy of cancer, especially appeared encoding a cytotoxin, for therapy of cancer, especially appeared is necessary for the unrestricted proliferative capacity of many human cancers. Mutation or dysregulation of the telomerase expression pathway may cause reactivation or dependences expression in cancer. Substances, cor uppequiation of telomerase expression in cancer. Substances, conception in the methods, can be used to block transcription from the TR gene promoter through interaction of the 5' regulatory sequences. These substances, e.g. antisense oligonucleotides, transcription factors, peptide nucleic acids and factors that disrupt signal transculction, are useful for cancer therapy. In particular, gene therapy vectors (specially pGT62-codAupp) comprising the promoter and a viral thymidine kinase gene can be used to convert a product of eggenially pGT62-codAupp) comprising the promoter and a viral thymidine coplasia can be used to convert a product, e.g. gancyclovir, so that neoplasia can be used to convert a product, e.g. gancyclovir, convert a product down regulation of telomerase RNA gene through manipulation of transcription factors may be effective anticancer therapy and the cloning of the hTR gene promoter allows the analysis of therapeutic molecules which modulate hTR promoter allows the analysis of therapeutic molecules which modulate hTR promoter active the proportion of the think the analysis of therapeutic molecules which modulate hTR promoter allows the analysis of therapeutic molecules which modulate hTR promoter active the promoter and the converted promoter allows the analysis of therapeutic molecules which modulate hTR promoter allows the analysis of therapeutic molecules which modulate hTR promoter allows the analysis of therapeutic molecules.
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                                                                                         Mouse and human telomerase RNA gene promoters, useful for tumor specific
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                                                                                                                                                                    invention relates to promoter regions from mouse and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
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with hTR promoter sequence element mutations
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Pred. No. 8.9;
0; Mismatches
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                                                                                                                                                                                                                                     Disclosure; Fig 19; 109pp; English
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WPI; 1999-479183/40.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sapiens.
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RESULT
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                     The invention relates to promoter regions from mouse and human telomerase
RNA (TR) component genes. The TR gene promoter can be linked to a
Reterologous gene, especially a gene encoding a cytotoxin, for therapy of
cancer, especially neoplasias. The telomerase is necessary for the
currestricted proliferative capacity of many human cancers. Mutation or
dysregulation of the telomerase repression pathway may cause reactivation
of upregulation of telomerase expression in cancer. Substances,
clentified in the methods, can be used to block transcription from the TR
gene promoter through interaction of the 5' regulatory sequences. These
substances, e.g. antisense oligonucleotides, transcription factors,
peptide nucleic acids and factors that disrupt signal transduction, are
useful for cancer therapy. In particular, gene therapy vectors
cuseful for cancer therapy. In particular, gene therapy vectors
cuseful for cancer therapy. In particular, gene therapy vectors
changes gene can be used to convert a products and a viral thymidine
kinase gene can be used to convert a producy, e.g. gancyclovir, so that
neoplasia can be controlled or treated. Direct down-regulation of
telomerase RNA gene through manipulation of transcription factors may be
effective anticancer therapy and the cloning of the hTR gene promoter
allows the analysis of therapeutic molecules which modulate hTR promoter
activity. Sequences AAZ07324-332 and AAZ00332-340 represent constructs
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                                                                                                                                                                                                                                                                                                                                                   Score 36.4; DB 1; Length 47; Pred. No. 17; 0; Mismatches 6; Indels
                                                                                                                                                                                                                                                                                                                                                                                                           6 GCGGAGGGTGGGCTGGGAGGGTGGTGGCCATTTTTTGTCTAACC 51
                                                                                                                                                                                                                                                                                                                                                                                                                                       2 drecadedrecereceraserasecraricerritriritricionale 47
                                                                                                                                                                                                                                                                                                                         Sequence 47 BP; 8 A; 7 C; 18 G; 14 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                            with hir promoter sequence element mutations
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Disclosure, Fig 19; 109pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAZ07296 standard; DNA; 38 BP
                                                                                                                                                                                                                                                                                                                                                  87.0%;
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                               Matches
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                                                                                                                                                     identified in the methods can be used to block transcription from the gene promoter through interaction of the 5' regulatory sequences. These substances, e.g. antisense oligonucleotides, transcription factors. peptide nucleic acids and fearors that disrupt signal transduction, useful for cancer therapy. In particular, gene therapy vectors (especially pGFG2-codAupp) comprising the promoter and a viral thymidine kinase gene can be used to convert a prodrug, e.g. gancyclovir, so that neoplasia can be used to convert a prodrug, e.g. gancyclovir, so that conparase RNA gene through manipulation of transcription factors may be effective anticancer therapy and the cloning of the NTR gene promoter allows the analysis of therapeutic molecules which modulate hTR promoter activity. Sequences AAZO7696-321 represent PCR primers used in cloning
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to promoter regions from mouse and human telomerase RNA (TR) component genes. The TR gene promoter can be linked to a heterologous gene, especially a gene encoding a cytotoxin, for therapy of cancer, especially neoplasias. The telomerase is necessary for the unrestricted proliferative capacity of many human cancers. Mutation or dysregulation of the telomerase repression pathway may cause reactivation or upregulation of telomerase expression in cancer. Substances, identified in the methods, can be used to block transcription from the TR
cancer, especially neoplasias. The telomerase is necessary for the unrestricted proliferative capacity of many human cancers. Mutation or dysregulation of the telomerase repression pathway may cause reactivation or upregulation of telomerase expression in cancer. Substances,
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14;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    and mutagenisis of human TR gene (hTR) promoter region
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 GGGTTGCGGAGGGTGGGCCTGGGAGGGGTGGTGGCC 36
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAZ07298 standard; DNA; 38 BP.
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Matches 36; Conservative
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Page 10

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                substances, e.g. antisense oligonucleotides, transcription factors, peptide nucleic acids and factors that disrupt signal transduction, are useful for cancer therapy. In particular, gene therapy vectors (especially pGT62-codAupp) comprising the promoter and a viral thymidine kinase gene can be used to convert a prodrug, e.g. gancyclovir, so that neoplasia can be controlled or treated. Direct down regulation of telomerase RNA gene through manipulation of transcription factors may be
                                                                                                                                                       effective anticancer therapy and the cloning of the hTR gene promoter allows the analysis of therapeutic molecules which modulate hTR promoter activity. Sequences AAZ07696-321 represent PCR primers used in cloning and mutagenisis of human TR gene (HTR) promoter region
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                                                                                                                                                                                                                                                                                                                                             Indels
promoter through interaction of the 5' regulatory
                                                                                                                                                                                                                                                             Sequence 38 BP; 3 A; 7 C; 21 G; 7 T; 0 U; 0 Other;
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Pred. No. 36;
0; Mismatches
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mutagenesis; PCR primer; ss.
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                                                                                                                                                                                                                                                                                                                      ilarity 91.7%;
Conservative
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Best Local Similarity
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kinase gene can be used to convert a prodrug, e.g. gancyclovir, so that neoplasia can be controlled or treated. Direct down-regulation of telomerase RNA gene through manipulation of transcription factors may be effective anticancer therapy and the cloning of the hTR gene promoter allows the analysis of therapeutic molecules which modulate hTR promoter activity. Sequences AA207669-321 represent PCR primers used in cloning and mutagenisis of human TR gene (hTR) promoter region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human, telomerase RNA component, anticancer therapy, purification, assay, vaccine, cancer, antisense oligonuclectide; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pruzan RA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present sequence represents an antisense oligonucleotide directed
                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Antisense oligonucleotide anti-P for human telomerase RNA component.
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                                                                                                                                                                 DB 1; Length 38;
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                                                                                                                                                                                                  Indels
                                                                                                                                 Sequence 38 BP; 5 A; 7 C; 20 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                 36
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                                                                                                                                                                                                                                 1 GGGTTGCCGAGGGTGGCCCTGGGAGGGGGTGGTGGCC
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                                                                                                                                                                Score 31.2; DE Pred. No. 36; 0; Mismatches
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"biotinylated"
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"biotinylated"
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                                                                                                                                                                 ch 6.9%;
1 Similarity 91.7%;
33; Conservative
                                                                                                                                                                                                                                                                                                                                                 AAV63645 standard; DNA; 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Atkinson EM,
                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
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/note=
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/note=
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                                                                                                                                                                 Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  modified base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (GERO-) GERON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                04-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               04-APR-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO9845450-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                                                                                                                                                                  AAV63645;
                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                    RESULT 14
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biochemical analysis of its activity, and in preparation of antibodies. Fragments of telomerase, or nucleic acid encoding them, are used in vaccines, and for treating over expression of telomerase, particularly in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New peptide nucleic acid (PNA) compounds that inhibit telomerase activity in mammalian cells is useful as probes to detect the RNA component of a mammalian telomerase.
                                                                                                                                                                                                                                                                                                                                                            Peptide nucleic acid; PNA; telomerase; ribonucleoprotein enzyme; cancer; inhibitor; neoplasia; neurodegenerative disease; aging; hyperplasia; AIDS; HIV; fungal infection; forensic identification; detect; tumour;
                                                                                                                          Gaps
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                                                                                            Similarity 100.0%; Pred. No. 28; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Corey
                                                                   Sequence 31 BP; 7 A; 9 C; 7 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Norton JC,
                                                                                                                                                   TIGICIAACCCTAACTGAGAAGGGCGTAGGC 72
                                                                                                                                                                              31 TTGTCTAACCCTAACTGAGAAGGGCGTAGGC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Wright WE, Piatyszek MA, Shay JW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 2; Col 37; 45pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              96US-00630019.
                                                                                                                                                                                                                                                                                                                                    Telomerase target sequence.
                                                                                                                                                                                                                                                  AAA37591 standard; RNA; 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (TEXA ) UNIV TEXAS SYSTEM
                                                                                                                                                                                                                                                                                                                                                                                                        paternity testing; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2000-292432/25.
                                                                                            Query Match
Best Local Simil
Matches 31; C
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                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
                                                                                                                                                                                                                                                                             AAA37591;
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                                                                                                                                                                                                                         RESULT 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to peptide nucleic acids (PNAs), comprising a sequence of 6-25 nucleobases, that inhibit telomerase activity in mammalian cells by hybridising to the RNA component of mammalian telomerase. The PNAs are useful as probes to detect the RNA component of
probes to detect or quantitate polynucleotides having a human telomerase RNA (hTR) sequence. PNA probes are also used for forensic identification of individuals, e.g. paternity testing, based on hTR gene restriction fragment length polymorphism (RFLP) pattern. PNAs are also useful as probes to detect the RNA component of a mammalian telomerase and as inhibitors of telomerase activity. The method of the present invention possibly at an earlier stage, before cells are detected as cancerous based on pathological characteristics. The diagnostic and prognostic methods of the present invention can be used to detect an immortal or neoplastic cell or tumour tissue or cancer of any origin, provided the cell expresses telomerase activity and its RNA component
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New peptide nucleic acids that hybridizes to the RNA component of mammalian telomerase, useful for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               6.9%; Score 31; DB 1; Length 31; 71.0%; Pred. No. 28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Indels
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/note= "Telomerase RNA active site"
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 31 BP; 8 A; 6 C; 8 G; 0 T; 9 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    40 TITIGICIAACCTAACTGAGAAGGGCGTAG 70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           9; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Corey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human telomerase RNA (hTR) strand #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 2; Col 37-38; 46pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
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97US-00838545.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14-FEB-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (TEXA ) UNIV TEXAS SYSTEM.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 71.0
Matches 22; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-638024/73.
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09-APR-1997;
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mammalian telomerase and as inhibitors of telomerase activity, or to detect and/or quantitate polynucleotide having the human telomerase RNA component (HTR) sequence, as well as in forensic identification of individuals, such as paternity teating or identification of criminal suspects or unknown descendants based on the hTR gene RFLP pattern. The PNAs an be further used for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases. The PNAs in combination with other pharmaceuticals (such as antimoplastic or rytostatic agents) can be used for treating neoplasia, hyperplasia, human immunodeficiency virus (HTV) infections, acquired immunodeficiency syndrome (AIDS) and associated pathologies, and other diseases characterised by abnormal telomere metabolism or telomerase activity. The present sequence repersenting hTR RNA strand #2 is used to test the inhibition of telomerase activity by the PNAs of the present
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                                                                                                                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human, Telomerase, vaccine, antibody, cancer; EF2H; nucleolin; antisense oligonucleotide; anti-P; ss.
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                                                                                                                                                                                                                                                            Score 31; DB 1; Length 31;
Pred. No. 28;
9; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Antisense oligonucleotide for human telomerase, anti-P.
                                                                                                                                                                                                                                    Sequence 31 BP; 8 A; 6 C; 8 G; 0 T; 9 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Lichtsteiner SP,
                                                                                                                                                                                                                                                                                                                      40 TITIGICIAACCCTAACTGAGAAGGGCGTAG 70
                                                                                                                                                                                                                                                                                                                                        UNUUGUCUAACCCUAACUGAGAAGGGCGUAG
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/note= "G is biotinylated"
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                                                                                                                                                                                                                                                            6.9%;
71.0%;
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                                                                                                                                                                                                                                                                               Similarity 71.0 22; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             nodified base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      04-AUG-1995;
04-APR-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
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                                                                                                                                                                                                            invention
                                                                                                                                                                                                                                                               Query Match
Best Local {
                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAS09472;
                                                                                                                                                                                                                                                                                             Matches
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Disclosure; Col 18; 29pp; English.

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purity compared with a crude extract of cells from adenovirus-transformed kidney cell line (293 cells) and when associated with telomerase RNA component has DNA polymerase activity and a molecular weight of 200-2000 kilo Daltons (kDa). The purified telomerase is useful for inducing a chamoral or cell-mediated immune response in an animal. Purified telomerase associated with over-expression of telomerase, such as cancer and for producing antibodies that recognize telomerase, which are useful as a ffinity agents in isolating the proteins and for detecting the presence of proteins in a sample, such as call or tissue. Identification confuencase and/or telomerase associated proteins are also useful for screening compounds to identify agents that alter the association of screening compounds to identify agents that alter the association of telomerase associated proteins are also useful for screening compounds to identify agents that alter the association of telomerase.
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                                                                                                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                                                                                                      6.9%; Score 31; DB 1; Length 31; 00.0%; Pred. No. 28;
                                                                                                                                                                                                                                                                                                                                                                          0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human telomerase antisense oligonucleotide primer AntiP.
                                                                                                                                                                                                                                                                                                  Sequence 31 BP; 7 A; 9 C; 7 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Telomerase; antisense; primer; AntiP; ss; cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                   42 TIGICIAACCCTAACTGAGAAGGGCGTAGGC 72
                                                                                                                                                                                                                                                                                                                                                          100.0%; Pred. No. 28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     31 TTGTCTAACCCTAACTGAGAAGGGCGTAGGC 1
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/note= "Biotinylated"
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/note= "Biotinylated"
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97US-00833377.
99US-00420056.
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                                                                                                                                                                                                                                                                                                                                                            Local Similarity 100. es 31; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          modified base
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04-APR-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABX10982;
                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABX10982/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            $$$$$$$$$$$$$$$$$$$$$$$$$
                                                                                                                                                                                                                                                                                                                                                                                                                                                         셤
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The sequence represents a biotinylated antisense oligonucleotide used in the purification of human telomerase. The invention relates to a purified human telomerase core enzyme protein comprising 2000-fold increased
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Purified human telomerase, useful for inducing immune response in animals, comprises several thousand folds increased purity compared with cytoplasmic crude cell preparations.
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measuring telomerase enzymatic activity of the composition in presence of a regulator. The telomerase protein of the invention may be used in developing and testing assays for measuring telomerase activity which are useful in characterising cancer and pre-cancer cells, for identifying and testing regulators of telomerase activity in in vitro assay and for preparing antibodies against telomerase. The mammalian telomerase protein of the invention is at least approximately 3000 fold more pure (in terms of telomerase activity per weight of protein) than a crude extract of cell from adenovirus- transformed kidney cell. Purified telomerase cativity bacomerase with the nayme's mechanism for developing mechanism- based regulators. The present sequence represents a human telomerase antisense oligonucleotide which has affinity to telomerase and is used to purify the telomerase protein of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention describes a method of obtaining mammalian telomerase protein (I). The method involves preparing enriched solution (ES) from a cell expressing telomerase where the component of (I) in ES is separated from other proteins expressed by cell by combining ES with oligonucleotide (O) having specific affinity for (I), and collecting protein bound to (O). The oligonucleotide comprises a retrievable label such as biotin and contains a sequence that is specifically recognized by telomerase protein. The oligonucleotide contains or does not contain the sequence (TTAGGG)3. The method further comprises combining a fraction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Obtaining telomerase, by preparing enriched solution from cell expressing telomerase, combining the solution with oligonucleotide having specific affinity for the protein and collecting protein bound to oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Weinrich SL, Atkinson EM, Lichtsteiner SP, Vasserot AP, Pruzan RA;
                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            mammalian telomerase protein; telomerase purification; telomere; anion exchange matrix; cation exchange matrix ; selectivity matrix; gel filtration chromatography; gradient centrifugation; antisense oligonucleotide; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human telomerase RNA component antisense oligonucleotide seq id
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                                                                                                                                                                                                                                                                                                                         Score 31; DB 1; Length 31;
Pred. No. 28;
                                                                                                                                                                                                                                                                                                                                                                 0; Indels
                                                                                                                                                                                                                                                                                    Sequence 31 BP; 7 A; 9 C; 7 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                      TTGTCTAACCCTAACTGAGAAGGGCGTAGGC 72
                                                                                                                                                                                                                                                                                                                                    100.0%; Pred. ....
                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; SEQ ID NO 1; 24pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ВР
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20-NOV-2000; 2000US-00717829.
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99US-00420056.
                                                                                                                                                                                                                                                                                                                            6.98;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 100.
Matches 31; Conservative
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containing telomerase protein with an anion exchange matrix, and collecting protein that binds the matrix, combining a fraction containing telomerase protein with a cation exchange matrix (such as a heparin matrix), and collecting protein that binds the matrix. The method comprises successively enriching fractions containing telomerase protein on several different ion exchange matrices and combining a fraction containing telomerase protein with an intermediate selectivity matrix, collecting protein that binds the matrix, where the intermediate selectivity matrix and separating a fraction containing the telomerase protein by gel filtration chromatography or gradient centrifugation. The telomerase is enriched from an extract of cells stably expressing telomerase. This sequence represents an antisense oligonucleotide to the RNA component of human telomerase that can be used in the purification
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                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Telomerase activity; therapy; cancer; cytostatic; antisense; ss.
                                                                                                                                                                                                                                                                                       ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Vasserot AP,
                                                                                                                                                                                                                                                        DB 1; Length 31;
28;
                                                                                                                                                                                                                                                                                     Indela
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human telomerase RNA antisense oligonucleotide, anti-P.
                                                                                                                                                                                                                            Sequence 31 BP; 7 A; 9 C; 7 G; 8 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                   42 TTGTCTAACCTAACTGAGAAGGGCGTAGGC 72
                                                                                                                                                                                                                                                                       100.0%; Pred. No. 28;
:ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                          TTGTCTAACCCTAACTGAGAAGGGCGTAGGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /*tag= b
/mod_base= OTHER
/note= "Biotin labelled"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /*tag= a
/mod_base= OTHER
/note= "Biotin labelled"
                                                                                                                                                                                                                                                           6.9%; Score 31;
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04-APR-1997; 97US-00833377.
18-0071-1997; 99US-00420056.
20-NOV-2000; 2000US-00717828
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                24-DEC-2002; 2002US-00330872
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    11-MAR-2004 (first entry)
                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 100.
Matches 31; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ᅼ
                                                                                                                                                                                                 nethod of the invention.
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ATKINSON E M.
LICHTSTEINER S
VASSEROT A P.
PRUZAN R A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            VPI; 2003-811733/76.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    modified base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      02-OCT-2003
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(VASS/)
(PRUZ/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (WEIN/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (ATKI/)
                                                                                                                                                                                                                                                                                                                                                                                             RESULT 20
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                            The invention relates to a method for identifying regulators of telomerase activity that may be useful for treating cancers. The method may be used to identify regulators e.g. antibodies, of telomerase activity which may be useful as cancer treatments. It has been found that found that the cells of many human cancers have telomerase activity. This helps explain why cancer cells continue dividing without becoming senescent. If telomerase activity in cancer cells can be inhibited, the cancer cells are expected to reach senescence and cease dividing. The present sequence is human telomerase antisense oligonulceotide used to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              mouse, rat and chinese hamster are all claimed. Antisense digonucleotides can be used to block the activity of the telomerase; probes and primers can be used in detection; vectors and host cells transformed with the isolated telomerase genes can be used for production of telomerases; RNA and DNA ribozymes and triplex forming oligonucleotides directed against the telomerase genes can be used therapeutically as can plasmids. A mouse which lacks the telomerase gene (also claimed) can be used for study of telomera regulation in vivo, and the role it plays in immortalisation. Three primers (AATI1040, AATI1043,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        - useful in studying
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The RNA components of (non) human mammalian telomerase(s) especially from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Telomerase; mammal; antisense; triplex forming oligonucleotide; plasmid; probe; primer; ribozyme; ss.
                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Marhuenda MA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Primer for production of telomerase antisense oligonucleotide.
                                                                                                                                                                                                                                           DB 1; Length 31; 28;
                                                                                                                                                                                                                                                                            0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RNA components of (non)human mammalian telomerase(s) cell senescence and immortalisation.
                                                                                                                                                                                                         Sequence 31 BP; 7 A; 9 C; 7 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                       42 TIGICIAACCTAACTGAGAAGGGCGTAGGC 72
                                                                                                                                                                                                                                                                                                                           31 TTGTCTAACCCTAACTGAGAAGGGCGTAGGC 1
                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Funk W,
                                                                                                                                                                           illustrate the method of the invention.
                                                                                                                                                                                                                                         6.9%; Score 31;
100.0%; Pred. No.
Disclosure; SEQ ID NO 1; 22pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Feng J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 1; Page 53; 85pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COLD SPRING HARBOR LAB
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95US-00387524.
95US-00485778.
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AAT11043 standard; DNA; 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                         Query Match 6.9
Best Local Similarity 100.
Matches 31; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1996-097428/10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GERON CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Villeponteau B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO9601614-A2
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13-FEB-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAT11043;
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AAT11044) were used to produce antisense oligonucleotides which were then
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           human - useful in identifying
                 used to produce antisense expression plasmids. AAT11040 was used alongside both AAT11043 and AAT11044 to produce two different antisense
                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RNA component; human; telomerase; lung fibroblast; cell line WI-38;
                                                                                                                                                                                                                                                                                                                                                                                                                                        RNA component of human telomerase antisense plasmid PCR primer G1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             recombinant production; synthesis; muťant; detection; mammalian; identification; modulating agent; neoplastic condition; transcriptional regulatory sequence; gene therapy; disease; polymerase chain reaction; antisense plasmid; PCR primer; ss.
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0
                                                                                                                         6.7%; Score 30; DB 1; Length 30;
100.0%; Pred. No. 32;
ive 0; Mismatches 0; Indels
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                                                                                      Sequence 30 BP; 13 A; 6 C; 11 G; 0 T; 0 U; 0 Other;
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                                                                                                                                                                                                77 IGCTTTTGCTCCCCCCCCCCTGTTTTCTC 106
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           candidate telomerase-modulating agents.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RNA component of mammalian telomerase,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 8; Page 80; 114pp; English.
                                                                                                                                                                                                                                                                                                                                 BP
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94US-00330123.
95US-00472802.
95US-00482115.
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                                                                                                      Query Match
Best Local Similarity 100.00
Conservative
                                                                                                                                                                                                                                                                                                                              AAT10298 standard; DNA; 30
                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
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Best Local Similarity
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                                                     molecules
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                                                                                                                                                                                                                                                  AAT10298;
                                                                                                                                                                                                                                                                                             RESULT 22
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Human; telomerase RNA component; anticancer therapy; purification; assay; vaccine; cancer; antisense oligonucleotide; ss.

/*tag= a /note= "biotinylated" Location/Qualifiers

Key modified_base

WO9845450-A1

15-OCT-1998

Homo sapiens

Synthetic.

97WO-US006012 97WO-US006012

04-APR-1997; 04-APR-1997;

Antisense oligonucleotide 14 for human telomerase RNA component.

(first entry)

15-FEB-1999

AAV63649;

AAV63649 standard; DNA; 30 BP.

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AAV63649/c
ID AAV63
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present sequence represents an antisense oligonucleotide directed against the human telomerase RNA component gene sequences. The oligonucleotide can be used as an affinity agent in the methods of the invention, which are used to purify human telomerase. The methods of the use of several sequential steps, including the use of two matrixes that bind molecules bearing negative charges, a matrix that binds molecules bearing negative charges, an matrix that binds molecules bearing negative charges, an affinity purification step and a size separation. Telomerase is a particular target of anticancer therapies, and is useful in assays for characterizing (pre) cancerous cells. Telomerase can also be used to screen for specific modulators, for biochemical analysis of its activity, and in preparation of antibodies. Fragments of telomerase, or nucleic acid encoding them, are used in vaccines, and for treating over expression of telomerase, particularly in
                                                                                                                                                                                                                                                                                                                                                                               Human; telomerase RNA component; anticancer therapy; purification; assay;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Atkinson EM, Lichtsteiner SP, Vasserot AP, Pruzan RA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Purification of telomerase on affinity material - useful for, e.g diagnosis and treatment of cancer.
                                                                                                                                                                                                                                                                                                                                   Antisense oligonucleotide 13 for human telomerase RNA component.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         6.7%; Score 30; DB 1; Length 30;
100.0%; Pred. No. 32;
tive 0; Mismatches 0; Indels
  0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 30 BP; 6 A; 6 C; 8 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                   vaccine; cancer; antisense oligonucleotide; ss.
                                              TGCTTTTGCTCCCCGCGCGCTGTTTTTCTC 106
Mismatches
                                                                       30 TGCTTTTGCTCCCGCGCGCGCTTTTTCTC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /*tag= a
/note= "biotinylated"
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                                                                                                                                                                                                 AAV63648 standard; DNA; 30
                                                                                                                                                                                                                                                                                     (first entry)
  30; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1998-594485/50.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Key
modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ното вартепв
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO9845450-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Kealey JT;
                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
                                                                                                                                                                                                                                             AAV63648;
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Best Local &
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Matches
  Matches
                                                                                                                                                      RESULT 23
AAV63648(XX
XX
AAV65
AAV66
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Pruzan RA;

Vasserot AP,

Lichtsteiner SP,

Atkinson EM,

Weinrich SL,

Kealey JT;

(GERO-) GERON

WPI; 1998-594485/50.

Purification of telomerase on affinity material - useful for, diagnosis and treatment of cancer.

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                                                      The present sequence represents an antisense oligonucleotide directed against the human telomerase RNA component gene sequences. The oligonucleotide can be used as an affinity agent in the methods of the invention, which are used to purify human telomerase. The methods of the the use of several sequential steps, including the use of two matrixes that bind molecules bearing negative charges, a matrix that binds molecules bearing negative charges, a matrix that binds molecules bearing positive charges, an affinity purification step and a size separation. Telomerase is a particular target of anticancer therapies, and is useful in assays for characterizing (pre) cancerous cells. Telomerase can also be used to screen for specific modulators, for biochemical analysis of its activity, and in preparation of antibodies. Fragments of telomerase, or nucleic acid encoding them, are used in vaccines, and for treating over expression of telomerase, particularly in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 30 BP; 8 A; 5 C; 13 G; 4 T; 0 U; 0 Other;
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100.0%; Pred. No. Jr.
7. 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     137 CCTGCCGCCTTCCACCGTTCATTCTAGAGC 166
Disclosure; Page 24; 76pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAV41175 standard; DNA; 30 BP
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AAV41175/C
ID AAV41
XX
AC AAV41
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AAACAAAAATGTCAGCTGCTGGCCGTTC 196

167

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AAACAAAAATGTCAGCTGCTGGCCCGTTC 1

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Sequences shown in AAV41169 to AAV41181 represent antisense
oligonucleotides to the RNA component of human telomerase (hTR). These
antisense oligonucleotides specifically hybridise to a nucleotide
sequence within an accessible region of the HTR, but that does not
bybridise to a sequence within the template region of hTR. These
component of human telomerase in a sample. This is useful for diagnosing
component of human telomerase in a sample. This is useful for diagnosing
cancer (especially neuroblastoma, bladder, colon and prostate cancer),
component of human telomerase and the inhibitory
can providing prognosis for a cancer patient. The inhibitory
coligonucleotides can inhibit the telomerase activity level in a cell by
interfering with transcription of the RNA component, decreasing the half-
coligonucleotides can inhibit the telomerase activity is inhibiting assembly of
the RNA component into the telomerase holoenzyme, or inhibiting the
polymerase activity of telomerase. These antisense oligonucleotides can
be used for inhibiting telomerase activity in both cultured cells and in
preventing cancer, for contraception or sterilisation, for
immunosuppression, and for selectively down-regulating specific branches
coliformatory thereapies or for treating infections by, e.g. yeast,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New polynucleotide(s) anti:sense to human telomerase - used for detecting or inhibiting human telomerase, e.g. for treating cancers, contraception, immuno-suppression or treating infection.
                                                                                           RNA component; human telomerase; antisense oligonucleotide; infection; neuroblastoma; bladder cancer; colon cancer; prostate cancer; cancer; contraception; sterilisation; immunosuppression; therapeutic; hTR; immune system down-regulation; anti-inflammatory therapy; ss.
                                                   RNA component of human telomerase (hTR) antisense oligo 21.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Kim NW, Wu F, Kealey JT, Pruzan R, Weinrich SL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 30 BP; 8 A; 5 C; 13 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               137 CCTGCCGCCTTCCACCGTTCATTCTAGAGC 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       6.7%; Score 30; DB
100.0%; Pred. No. 32;
ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CCTGCCGCCTTCCACCGTTCATTCTAGAGC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 11; Page 65; 80pp; English
                                                                                                                                                                                                                                                                                                                           97WO-US023619.
                                                                                                                                                                                                                                                                                                                                                             96US-00770564.
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                     (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1998-377670/32.
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                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                                WO9828442-A1
                                                                                                                                                                                                                                                                                                                       19-DEC-1997;
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                   08-OCT-1998
                                                                                                                                                                                                                                                                                                                                                               20-DEC-1996;
                                                                                                                                                                                                                                                                                   02-JUL-1998.
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                                                                                                                                                                                          Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
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ID AAV4
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AC AAV4
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Sequences shown in AAV41169 to AAV41181 represent antisense
oligonucleotides to the RNA component of human telomerase (hTR). These
antisense oligonucleotides specifically hybridise to a nucleotide
cequence within an accessible region of the hTR, but that does not
bybridise to a sequence within the template region of hTR. These
coligonucleotides may specifically be used for detection of an RNA
component of human telomerase in a sample. This is useful for diagnosing
cancer (especially neuroblastoma, bladder, colon and prostate cancer),
and providing prognosis for a cancer patient. The inhibitory
cancer (especially neuroblastoma, bladder, colon and prostate cancer),
and providing prognosis for a cancer patient. The inhibitory
clife of the telomerase RNA component transcript, inhibiting assembly of
interfering with transcription of the RNA component, decreasing the half-
clife of the telomerase RNA component transcript, inhibiting assembly of
the RNA component into the telomerase holoenzyme, or inhibiting the
polymerase activity of telomerase activity in both cultured cells and in
be used for inhibiting telomerase activity in both cultured cells and in
preventing cancer, for contraception or sterilisation, for
immunosuppression, and for selectively down-regulating specific branches
cof the immune system, e.g. a specific subset of T-cells, in anti-
inflammatory therapies or for treating infections by, e.g. yeast,
clife of the fulliphical person of the treating infections by, e.g. yeast,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New polynucleotide(s) anti:sense to human telomerase - used for detecting or inhibiting human telomerase, e.g. for treating cancers, contraception, immuno-suppression or treating infection.
                                                                             RNA component; human telomerase; antisense oligonucleotide; infection; neuroblastoma; bladder cancer; colon cancer; prostate cancer; cancer; contraception; sterilisation; immunosuppression; therapeutic; hTR; immune system down-regulation; anti-inflammatory therapy; ss.
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0
                                        RNA component of human telomerase (hTR) antisense oligo 16.
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100.0%; Pred. No. 32;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pruzan R, Weinrich SL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 30 BP; 6 A; 10 C; 9 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        290 CTGCCACCGCGAAGAGTTGGGCTCTGTCAG 319
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 11; Page 65; 80pp; English.
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96US-00770565.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Kim NW, Wu F, Kealey JT,
08-OCT-1998 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                               WO9828442-A1.
                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                19-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                       20-DEC-1996;
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Best Local Simil
Matches 30; C
                                                                                                                                                                                                                                                                                     02-JUL-1998
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                                                                                                                                                                                  Synthetic
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ID AAZ2:
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Gaps ;

1; Length 30; 0; Indels

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This invention describes a novel composition comprising human telomerase having at least 2000-fold (preferably at least 6000-fold) increased relative purity compared with crude extract of cells from adenovirus transformed kidney cell line. The composition is useful for eliciting an immune response in animals and may therefore be used as a vaccine for treating diseases associated with the overexpression of telomerase e.g. cancer. AA223626-Z23637 represent oligomucleotides used in the isolation of human clone 28-1 which contains a fragment of the human telomerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               human; immune response; cancer; vaccine; treatment; disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
6.7%; Score 30; DB 1; Length 30;
Best Local Similarity 100.0%; Pred. No. 32;
Matches 30; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                           Compositions comprising human telomerase, useful for tree associated with overexpression of telomerase e.g. cancer.
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/note= "5'-biotinylated guanosine"
 /*tag= a
/note= "5'-biotinylated guanosine"
                                                                                                                                                                                                              Weinrich SL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                167 AAACAAAAATGTCAGCTGCTGGCCGGTTC 196
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30 AAACAAAAATGTCAGCTGCTGGCCGTTC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                described in the method of the invention
                                                                                                                                                                                                                                                                                                                                                Disclosure; Col 45-46; 34pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                              Lichtsteiner SP,
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ID AAZ23631 standard; DNA; 30
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/*tag=
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                                                                                                                                                                                CORP
                                                                                                                                                                                (GERO-) GERON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Key
modified_base
                                                                                                             04-APR-1997;
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                                                                                                                                                                                                             Atkinson EM,
Vasserot AP;
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                                              US5968506-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US5968506-A
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                                                                                19-0CT-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAZ23631;
                                                                                                                                                                                                                                 Jasserot
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  원
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ò
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This invention describes a novel composition comprising human telomerase relative purity compared with created for following at least 6000-601d) increased relative purity compared with compession of cells from adenovirus-transformed kidney cell line. The composition is useful for eliciting an immune response in animals and may therefore be used as a vaccine for treating diseases associated with the overexpression of telomerase e.g. cancer. AAZ23626-Z23637 represent oligomucleotides used in the isolation of human clone 28-1 which contains a fragment of the human telomerase
                                               Telomerase; human; immune response; cancer; vaccine; treatment; disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 human; immune response; cancer; vaccine; treatment; disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Compositions comprising human telomerase, useful for treating diseases associated with overexpression of telomerase e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                Lichtsteiner SP, Weinrich SL, Pruzan RA, Kealey JT;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human clone 28-1 telomerase oligonucleotide oligo-13.
                 Human clone 28-1 telomerase oligonucleotide anti-P.
                                                                                                                                                                                /*tag= a
/note= "5'-biotinylated guanosine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 30 BP; 6 A; 9 C; 7 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      6.7%; Sco...
100.0%; Pred. No. ...
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Col 43-44; 34pp; English.
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                                                                                                                                               Location/Qualifiers
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                                                                                                                                                                                                                                                                                               97US-00833377
                                                                                                                                                                                                                                                                                                                               95US-00510736
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Best Local Similarity 100.
Matches 30; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1999-590379/50.
                                                                                                                                               Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                  (GERO-) GERON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  modified base
                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                 04-APR-1997;
                                                                                                                                                                                                                                                                                                                                 04-AUG-1995;
                                                                                                                                                                                                                                                                                                                                                                                                Atkinson EM,
Vasserot AP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    07-JAN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Telomerase;
                                                                                                                                                                                                                                USS968506-A
                                                                                                                                                                                                                                                                19-0CT-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    primer; BB
                                                                primer; ss
                                                                                               Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic
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Gaps

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useful for treating diseases

Kealey JT;

Pruzan RA,

US09436060A-16.rng.sl

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The invention relates to polynucleotide inhibitors (I) and methods for inhibiting telomerase activity. (I) are useful in inhibiting telomerase activity. (I) are useful in inhibiting telomerase activity and proliferation of a telomerase positive cell, and in manufacturing a medicament for inhibiting telomerase activity in a cell and in treating a medicament for inhibiting telomerase activity in a cell and in treating a tumour or in manufacturing a medicament for the Cueful in treating a tumour or in manufacturing a medicament for the cueful in treating a tumour or in manufacturing a medicament for the cueful in treating a tumour or in manufacturing a medicament for the cueful in treating a tumour or in manufacturing a medicament for the cueful in treating to detecting RNA or DNA. Inhibition of telomerase activity in cells in vivo is useful in prophylactic and therapeutic methods of treating encer and other disorders involving inappropriate expression of telomerase, and in treating veterinary proliferative chases. Inhibition of telomerase in hammatopoietic stem cells is useful continued to the immune of the continued that in the cells is useful continued to the continued that the continued tha
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              branches of \dot{L}he immune system. The present sequence represents human telomerase polynucleotide inhibitor #9, as described in the method of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                useful for inducing immune response in thousand folds increased purity compared with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human, Telomerase, vaccine, antibody, cancer; EF2H, nucleolin; antisense oligonucleotide; Oligo 14; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Vasserot AP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       6.7%; Score 30; DB 1; Length 30;
100.0%; Pred. No. 32;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Antisense oligonucleotide for human telomerase, Oligo 14.
tumors, Hodgkin's disease, or inflammatory conditions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 30 BP; 8 A; 5 C; 13 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Lichtsteiner SP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                137 CCTGCCGCCTTCCACCGTTCATCTAGAGC 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /mod_base= G
/note= "G is biotinylated"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30 CCTGCCGCCTTCCACCGTTCATTCTAGAGC
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                                                                    3; Page 32; 48pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAS09476 standard; DNA; 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Weinrich SL, Atkinson EM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
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Best Local Similarity 100.
Matches 30; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAS09476;
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AAS09476/c
THE STATE OF THE S
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                                                                                                                                                                                                                                                                                                                                                                                                                                  This invention describes a novel composition comprising human telomerase having at least 2000-fold (preferably at least 6000-fold) increased relative purity compared with crude extract of cells from adenovirus-transformed kidney cell line. The composition is useful for eliciting an immune response in animals and may therefore be used as a vaccine for treating diseases associated with the overexpression of telomerase e.g. cancer. AAZ213626-Z23637 represent oligonucleotides used in the isolation of human clone 28-1 which contains a fragment of the human telomerase described in the method of the invention
                                                                                                                                                                                                                                                       Compositions comprising human telomerase, useful for treating diseases associated with overexpression of telomerase e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; telomerase; hTR; cytostatic; anti-inflammatory; adenocarcinoma;
breast; prostate; colon; mixed cell leukaemia; Hodgkin's disease;
fertility; inflammatory condition; tumour; cancer; veterinary;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                        Kealey JT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             .
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                                                                        Pruzan RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 30; DB 1; Length 30;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 30 BP; 8 A; 5 C; 13 G; 4 T; 0 U; 0 Other;
                                                                        Lichtsteiner SP, Weinrich SL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human telomerase polynucleotide inhibitor #9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         immunosuppression; telomerase inhibitor; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 137 CCTGCCGCCTTCCACCGTTCATTCTAGAGC 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Pred. No. 34, tive 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Col 45-46; 34pp; English.
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/mod base=
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(GERO-) GERON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (GERO-) GERON CORP.
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modified_base
                                                                        AP;
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                                                                        Atkinson
                                                                                                              Vasserot
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ID AAS159;
XA AC AAS159;
XX AC AAS159;
XX Human |
XX Human |
XX Homo 8
XX Homo 9
XX Homo 9
XX Homo 9
XX Homo 11-OCT
XX Homo 1
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Gaps

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Pruzan RA;

13

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The sequence represents a biotinylated antisense oligonucleotide used in the purification of human telomerase. The invention relates to a purified human telomerase core enzyme protein comprising 2000-fold increased purity compared with a crude extract of cells from adenovirus-transformed purity compared with a crude extract of cells from adenovirus-transformed kidney cell line (293 cells) and when associated with telomerase RNA component has DNA polymerase activity and a molecular weight of 200-2000 killo Daltons (kDa). The purified telomerase is useful for inducing a humoral or cell-mediated immune response in an animal. Purified telomerase or immunogenic fragments are useful as vaccines for treating diseases associated with over-expression of telomerase, such as cancer and for producing antibodies that recognize telomerase, which are useful as affinity agents in isolating the proteins and for detecting the presence of proteins in a sample, such as cell or tissue. Identification of telomerase and/or telomerase associated proteins are also useful for carcening compounds to identify agents that alter the association of telomerase associated proteins are also useful for carcening compounds to identify agents that alter the association of telomerase associated proteins are also useful for carcening compounds to identify agents that alter the association of telomerase.
                                                                                                                                                                                                                                                                                                                                                   Sequence 30 BP; 6 A; 6 C; 8 G; 10 T; 0 U; 0 Other;
               Disclosure, Col 18; 29pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
nes 30; Conserv
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                    Query Match
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ABA91517/c
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                                                           The sequence represents a biotinylated antisense oligonucleotide used in the purification of human telomerase. The invention relates to a purified human telomerase core enzyme protein comprising 2000-fold increased purity compared with a crude extract of calls from adenovirus-transformed purity compared with a crude extract of calls from adenovirus-transformed component has DNA polymerase activity and a molecular weight of 200-2000 kilo baltons (kDa). The purified telomerase is useful for inducing a humoral or cell-mediated immune response in an animal. Purified telomerase or immunogenic fragments are useful as vaccines for treating diseases associated with over-expression of telomerase, which are useful as a fifnity agents in isolating the proteins and for detecting the proteins of proteins in a sample, such as cell or tissue. Identification of telomerase aids in disgnosis of cancer or pre-cancerous states.

Telomerase and/or telomerase associated proteins are also useful for screening compounds to identify agents that alter the association of telomerase and proteins, such as nucleolin or EF2H with telomerase
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Purified human telomerase, useful for inducing immune response in animals, comprises several thousand folds increased purity compared with cytoplasmic crude cell preparations.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pruzan RA;
                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Antisense oligonucleotide for human telomerase, Oligo 13.
                                                                                                                                                                                                                                                                                                                                                                                                     Score 30; DB 1; Length 30;
                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Indels
                                                                                                                                                                                                                                                                                                                                                                    Sequence 30 BP; 8 A; 5 C; 13 G; 4 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                            6.7%; Score ...
100.0%; Pred. No. 32;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     137 CCTGCCGCCTTCCACCGTTCTTCTAGAGC 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note= "G is biotinylated"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      30 cerececerrecaceerrearreragase 1
cytoplasmic crude cell preparations.
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                                 Disclosure; Col 18; 29pp; English
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AAS09475 standard; DNA; 30 BP.
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mod base= G
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97US-00833377
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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Best Local Similarity 100.
Matches 30; Conservative
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Kealey JT;
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04-APR-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       Silver staining, signal detection; nanoparticle; diagnosis; screening;
                                                 Gaps
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DB 1; Length 30; 32;
                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                          Oligonucleotide used in gold nanoparticle complex.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /*tag= a
/note= "phosphoramidate linkage"
                                                                                              167 AAACAAAAATGTCAGCTGCTGGCCCGTTC 196
              100.0%; Pred. ..
                                                                                                                                           AAACAAAAATGTCAGCTGCTGGCCCGTTC 1
  6.7%; Score 30;
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                                                                                                                                                                                                                                                                  ABA91517 standard; DNA; 30 BP
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                                               Conservative
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The present sequence is that of an oligonuclectide prepared using phosphoramidite chemistry. A nanoparticle conjugate was prepared by joining the 5'-mercaptoalkyl oligonuclectide to gold nanoparticles. The nanoparticle conjugate was lided to enhance the sandwich amplification of a silver signal on glass slides containing silver staining. This illustrated the method of the invention, which is based on the discoveries that: (1) gold nanoparticles coated with oligonuclectides bind to silver that has previously been deposited on gold nanoparticle-oligonuclectide conjugates immobilised by hybridisation on a glass substrate or plate; and (2) that the gold-nanoparticle-oligonuclectide-silver-gold oligonuclectide conjugates immobilised by hybridisation on a glass substrate or plate; and (2) that the discoveries were applied to a method for amplifying signal by chancing silver deposition in detecting systems where the formation of silver spot serves as a reporter for the presence of a molecule.

Companies and assays where the target molecules is bound to a substrate or is captured by a capture molecule is bound to a tissue sample) and assays where the target molecule is bound to a cutility in increasing the signal strength in diagnostic and screening applications involving detection of target molecules arrayed at discrete positions on a solid surface. It provides a means for greatly enhancing the sensitivity of tests carried out on microarrays or microchips. The method is simple, economical, and provides a large enhancement in signal
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human telomerase antisense oligonucleotide primer oligol3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     6.7%; Score 30; DB 1; Length 30;
100.0%; Pred. No. 32;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Telomerase; antisense; primer; oligol3; ss; cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 30 BP; 8 A; 5 C; 13 G; 4 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CCTGCCGCCTTCCACCGTTCATTCTAGAGC 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CCTGCCGCCTTCCACCGTTCATTCTAGAGC 1
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/note= "Biotinylated"
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97US-00833377.
99US-00420056.
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ABX10985 standard; DNA; 30
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Best Local Similarity 100.
Matches 30; Conservative
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04-APR-1997;
18-OCT-1999;
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This invention relates to a purified human telomerase protein, which when associated with telomerase RNA component has DNA polymerase activity.

Also disclosed in the specification is a method for assessing a regulator (preferably a telomerase inhibitor or activator of telomerase involves measuring telomerase enzymatic activity of the composition in presence of a regulator. The telomerase protein of the invention may be used in the developing and testing assays for measuring telomerase activity which are useful in characterising cancer and pre-cancer cells, for identifying and testing antibodies against telomerase. The mammalian telomerase protein of the invention is at least approximately 3000 fold more pure (in terms of telomerase activity per weight of protein) than a crude extract of cell from adenovirus- transformed kidney cell. Purified telomerase (cell from adenovirus- transformed kidney cell. Purified telomerase for feacilitates a thorough biochemical analysis of the enzyme's mechanism for developing mechanism based regulators. The present sequence represents a chuman telomerase antisense oligonucleotide which has affinity to telomerase and is used to purify the telomerase protein of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human telomerase antisense oligonucleotide primer oligol4.
                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1; Length 30;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
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                                                                                                                                                                                                                                                                                                                                                                                   Sequence 30 BP; 6 A; 6 C; 8 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
telomerase protein with telomerase RNA component
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                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; Pred. No. 32; tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         167 AAACAAAAATGTCAGCTGCTGGCCCGTTC 196
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          30 AAACAAAAATGTCAGCTGCTGGCCCGTTC 1
                                                                                                                                                                                                                                                                                                                                                                                                                      Score 30;
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/note= "Biotinylated"
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                                    Disclosure; Col 9; 24pp; English.
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97US-00833377.
99US-00420056.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABX10986 standard; DNA; 30
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                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
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modified_base
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04-APR-1997;
18-OCT-1999;
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ABX10986/c
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Also disclosed with telomerase RNA component has DNA polymerase activity.

Also disclosed in the specification is a method for assessing a regulator (preferably a telomerase RNA component has DNA polymerase activity.

Also disclosed in the specification is a method for assessing a regulator of preferably a telomerase involves measuring telomerase enzymatic activity of the composition in presence of a regulator. The telomerase protein of the invention may be used in developing and testing assays for measuring telomerase activity which are useful in characterising cancer and pre-cancer cells, for identifying and testing regulators of telomerase activity in in vitro assay and for preparing antibodies against telomerase. The mammalian telomerase protein of the invention is at least approximately 3000 fold more pure (in terms of telomerase activity per weight of protein) than a crude extract of cell from adenovirus- transformed kidney cell. Purified telomerase (activity per weight of protein) than a crude extract of facilitates a thorough biochemical analysis of the enzyme's mechanism for developing mechanism- based regulators. The present sequence represents a tunnary endeman and affinity to buman telomerase and sequence of the enzyme's mechanism for the numan telomerase antisense oligonucleotide which has affinity to
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invention relates to a purified human telomerase protein, which when
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                                                                                                                                                                                                                                                                                                                                              telomerase and is used to purify the telomerase protein of the invention
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97US-00833377.
99US-00420056.
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protein bound to (0). The oligonucleotide comprises a retrievable label such as biotin and contains a sequence that is specifically recognized by telometrase protein. The oligonucleotide contains or does not contain the sequence (TTAGGG): The method further comprises combining a fraction containing telomerase protein with an anion exchange matrix, and collecting protein that binds the matrix, combining a fraction containing telomerase protein with a cation exchange matrix. The method telomerase protein that binds the matrix. The method comprises successively enriching fractions containing telomerase protein containing telomerase protein on several different ion exchange matrices and combining a fraction on several different ion exchange matrix, where the intermediate containing telomerase protein with an intermediate selectivity matrix and separating a fraction containing the telomerase protein that binds the matrix, where the intermediate selectivity matrix and separating a fraction containing the telomerase protein that binds a fraction containing the telomerase protein that binds a fraction containing the telomerase protein that binds and a fraction containing the telomerase protein that binds and separating a fraction containing the telomerase protein that sequence represents an an extract of cells stably expressing telomerase. This sequence represents an an extract of cells achieved the tothe sequence represents an an extract of cells achieved the tothe sequence represents an an extract of cells achieved the tothe sequence represents an an extract of cells achieved to the sequence represents an an extract of cells achieved the tothe sequence represents an an extract of cells achieved the tothe sequence represents an an extract of cells achieved the tothe partitication that the partitication can be used in the purfit of the sequence and an extract of cells achieved the sequence represents and the telement of the sequence of the 
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04-APR-1997;
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Best Loca
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        protein bound to (0). The oligonaclectide comprises a retrievable label such as biotin and contains a sequence that is specifically recognized by telemerase protein. The oligonaclectide contains or does not contain the sequence (TTMGGG). The method further comprises combining a fraction containing telomerase protein with an anion exchange matrix, and collecting protein that binds the matrix, combining a fraction containing telomerase protein that binds the matrix. The method containing protein that binds the matrix. The method containing telomerase protein that binds the matrix. The method of matrix), and collecting protein that binds the matrix. The method containing telomerase protein with an intermediate selectivity matrix, on several different ion exchange matrices and combining a fraction on several different ion exchange matrix, where the intermediate collecting protein that binds the matrix, where the intermediate selectivity matrix, and separating a fraction containing the telomerase protein with an intermediate selectivity matrix and separating a fraction containing the telomerase protein chromatography or gradient centrifugation. The protein by gel filtration chromatography or gradient centrifugation. The collecting protein the matrix of cells stably expressing the matrix of the matrix of the centrifugation. The collection process is protein the centrifugation.
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                                                                                                                                                                                                                                                        telomerase. This sequence represents an antisense oligonucleotide to the RNA component of human telomerase that can be used in the purification
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 oligonucleotide (0) having specific affinity for (I),
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100.0%; Pred. No. 5.,
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/mod_base= OTHER
/note= "Biotin labelled"
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04-ARE-1997; 97US-0083377.
18-OCT-1999; 99US-00420056.
20-NOV-2000; 2000US-00717828
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ATKINSON E M.
LICHTSTEINER S
VASSEROT A P.
PRUZAN R A.
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Best Local Similarity
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(PRUZ/)
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telomerase activity that may be useful for treating cancers. The method may be used to identify regulators e.g. antibodies, of telomerase activity which may be useful as cancer treatments. It has been found that found that the cells of many human cancers have telomerase activity. This helps explain why cancer cells continue dividing without becoming senescent. If telomerase activity in cancer cells can be inhibited, the cancer cells are expected to reach senescence and cease dividing. The present sequence is human telomerase antisense oligonulceotide used to illustrate the method of the invention.
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                              The invention relates to a method for identifying regulators of
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                                                                                                                                                                                                                                                         6.7%; Score 30; DB 1; Length 30; 100.0%; Pred. No. 32;
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'''A 0; Mismatches
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Disclosure; SEQ ID NO 4; 22pp; English.
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04-APR-1997; 97US-0083377.
18-OCT-1999; 99US-00420056.
20-NOV-2000; 2000US-00718286.
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ID ADG62874 standard; DNA; 30
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ATKINSON E M.
LICHTSTEINER S P
VASSEROT A P.
PRUZAN R A.
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                                                                                                                                                                                                                                                         Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (ATK1/)
(LICH/)
(VASS/)
(PRUZ/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Identifying telomerase regulators useful for treating cancer.
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US09436060A-16.rng.sl

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present sequence represents an antisense oligonucleotide directed against the human telomerase RNA component gene sequences. The oligonucleotide can be used as an affinity agent in the methods of the invention, which were used to purify human telomerase. The methods of the the use of several sequential steps, including the use of two matrixes that bind molecules bearing negative charges, a matrix that binds molecules bearing negative charges, a matrix that binds molecules bearing positive charges, an affinity purification step and a size separation. Telomerase is a particular target of anticancer therapies, and is useful in assays for characterizing (pre) cancerous cells. Telomerase can also be used to screen for specific modulators, for biochemical analysis of its activity, and in preparation of antibodies. Fragments of telomerase, or nucleic acid encoding them, are used in vaccines, and for treating over expression of telomerase, particularly in
activity which may be useful as cancer treatments. It has been found that found that the cells of many human cancers have telomerase activity. This helps explain why cancer cells continue dividing without becoming senescent. If telomerase activity in cancer cells can be inhibited, the cancer cells are expected to reads senescence and cease dividing. The present sequence is human telomerase antisense oligonulceotide used to illustrate the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                     Human; telomerase RNA component; anticancer therapy; purification; assay; vaccine; cancer; antisense oligonucleotide; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pruzan RA;
                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Purification of telomerase on affinity material - useful for, e.g.
diagnosis and treatment of cancer.
                                                                                                                                                           ô
                                                                                                                                                                                                                                                                                                                                                             Antisense oligonucleotide 5 for human telomerase RNA component.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Vasserot AP,
                                                                                                                                 Query Match 6.7%; Score 30; DB 1; Length 30; Best Local Similarity 100.0%; Pred. No. 32; Matches 30; Conservative 0; Mismatches 0; Indels
                                                                                                        Sequence 30 BP; 8 A; 5 C; 13 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Lichtsteiner SP,
                                                                                                                                                                                     137 CCTGCCGCCTTCCACCGTTCTAGAGC 166
                                                                                                                                                                                                              ccreccerrecaccerrearreragase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /*tag= a
/note= "biotinylated"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 3; Page 47; 76pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           97WO-US006012
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    97WO-US006012
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Atkinson EM,
                                                                                                                                                                                                                                                                       647/c
AAV63647 standard; DNA; 30
                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1998-594485/50.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (GERO-) GERON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Weinrich SL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  04-APR-1997;
                                                                                                                                                                                                                                                                                                                                     15-FEB-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15-0CT-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Kealey JT;
                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
                                                                                                                                                                                                               30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This invention describes a novel composition comprising human telomerase having at least 2000-fold (preferably at least 6000-fold) increased relative purity compared with crude extract of cells from adenovirus-transformed kidney cell line. The composition is useful for eliciting an immune response in animals and may therefore be used as a vaccine for cracting diseases associated with the overexpression of telomerase e.g. cancer. AAZ23656-Z23637 represent oligomotoleotides used in the isolation of human clone 20-1 which contains a fragment of the human telomerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               human; immune response; cancer; vaccine; treatment; disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Compositions comprising human telomerase, useful for treating diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gapa
                                                                                                                                Gape
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                                                                                   DB 1; Length 30;
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                                                                                                                             Indele
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              associated with overexpression of telomerase e.g. cancer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pruzan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human clone 28-1 telomerase oligonucleotide oligo-5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /*tag= a
/note= "5'-biotinylated guanosine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 30 BP; 4 A; 11 C; 9 G; 6 T; 0 U; 0 Other;
                                       Seguence 30 BP; 4 A; 11 C; 9 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Weinrich SL,
                                                                               Score 28.4; DB Pred. No. 45; 0; Mismatches
                                                                                                                                                                       412 GAGCTGTGGGACGTGCACCCAGGACTCGGC 441
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GAGCTGTGGGACGTGCACCCAGGACTCGGC 441
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 28.4;
Pred. No. 45;
                                                                                                                                                                                                                   GAGCTATGGGACGTGCACCCAGGACTCGGC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30 GAGCTATGGGACGTGCACCCAGGACTCGGC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     described in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Col 43-44; 34pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Lichtsteiner SP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       97US-00833377.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    95US-00510736
                                                                                   6.3%;
                                                                                                                                                                                                                                                                                                                                 AAZ23629 standard; DNA; 30
                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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Best Local Similarity
Matches 29; Conserv
                                                                        Query Match
Best Local Similarity
Matches 29; Conserv
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     04-APR-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EM,
AP;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Telomerase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US5968506-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19-OCT-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                                             AAZ23629;
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cancer
                                                                                                                                                                                                                                                                                         RESULT 41
                                                                                                                                                                                                                                                                                                              AAZ23629,
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US09436060A-16.rng.sl

Human telomerase antisense oligonucleotide primer oligos.

(first entry)

17-AUG-2003

ABX10984;

ABX10984 standard; DNA; 30 BP.

RESULT 43 ABX10984/c

Pelomerase; antisense; primer; oligo5; ss; cancer.

Location/Qualifiers

Key modified_base

Synthetic

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The sequence represents a biotinylated antisense oligonucleotide used in the purification of human telomerase. The invention relates to a purified human telomerase core enzyme protein comprising 2000-fold increased purity compared with a crude extract of cells from adenovirus-transformed kidney cell line (293 cells) and when associated with telomerase RNA component has DNA polymerase activity and a molecular weight of 200-2000 kilo Daltons (kDa). The purified telomerase is useful for inducing a humoral or cell-mediated immune response in an animal. Purified telomerase or immunogenic fragments are useful as vaccines for treating diseases associated with over-expression of telomerase, such as cancer and for producing antibodies that recognize telomerase, which are useful as affinity agents in isolating the proteins and for detecting the presence of proteins in a sample, such as cell or tissue. Identification of telomerase and/or telomerase associated proteins are also useful for screening compounds to identify agents that alter the association of telomerase associated proteins are also useful for screening compounds to identify agents that alter the association of telomerase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Purified human telomerase, useful for inducing immune response in animals, comprises several thousand folds increased purity compared with cytoplasmic crude cell preparations.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Atkinson EM, Lichtsteiner SP, Vasserot AP, Pruzan RA;
                                                                                                                                                                            Human, Telomerase, vaccine, antibody, cancer; BF2H; nucleolin; antisense oligonucleotide; Oligo 5; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              6.3%; Score 28.4; DB 1; Length 30; 96.7%; Pred. No. 45; tive 0; Mismatches 1; Indels
                                                                                                                                           Antisense oligonucleotide for human telomerase, Oligo 5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 30 BP; 4 A; 11 C; 9 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                        /*tag= a
/mod_base= G
/note= "G is biotinylated"
                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Col 18; 29pp; English.
                                  AAS09474 standard; DNA; 30 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      99US-00420056
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        95US-00510736
97US-00833377
                                                                                                       (first entry)
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                                                                                                                                                                                                                                                                      Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                US6261556-B1.
                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                      18-OCT-1999;
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04-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Weinrich SL,
                                                                                                                                                                                                                                                                                                                                                                                                                    17-JUL-2001.
                                                                                                       24-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Kealey JT;
                                                                    AAS09474;
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             RESULT 42
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Also disclosed in the specification is a method for assessing a regulator. Also disclosed in the specification is a method for assessing a regulator. Also disclosed in the specification is a method for assessing a regulator or ellomerase inhibitor or activator of telomerase involves measuring telomerase enzymatic activity of the composition in presence of a regulator. The telomerase protein of the invention may be used in developing and testing assays for measuring telomerase activity which are useful in characterising cancer and pre-cancer cells, for identifying and testing regulators of telomerase activity in in vitro assay and for preparing antibodies against telomerase. The mammalian telomerase protein of telomerase activity per weight of protein) than a crude extract of cell from adenovirus- transformed kidney cell. Purified telomerase (a facilitates a thorough biochemical analysis of the enzyme's mechanism for developing mechanism- based regulators. The present sequence represents a chuman telomerase antisense oligonuclocide which has affinity to thome telomerase and is used to purify the telomerase protein of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This invention relates to a purified human telomerase protein, which when
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pruzan RA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Composition useful e.g. in diagnosis of cancer comprises complex of telomerase protein with telomerase RNA component.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 .;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             6.3%; Score 28.4; DB 1; Length 30; 96.7%; Pred. No. 45;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 30 BP; 4 A; 11 C; 9 G; 6 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
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                                                                                                                                                                                                         /*tag= a
/note= "Biotinylated"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 13; Col 9; 24pp; English.
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97US-00833377.
99US-00420056.
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ID ADC35650 standard; DNA; 30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-465598/44.
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hes 29; Conserv
                                                                                                                                                                                                                                                                                                                                                                04-AUG-1995;
                                                                                                                                                                                                                                                            US6517834-B1
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                                                                                                                                                                                                                                                                                               11-FEB-2003
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Matches
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8
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ö

Gaps

; 0

412 GAGCTGTGGGACGTGCACCCAGGACTCGGC 441

Best Local Similarity 96.7 Matches 29, Conservative

GAGCTATGGGACGTGCACCCAGGACTCGGC 1

30

à 셤 Telomerase activity; therapy; cancer; cytostatic; antisense; ss.

/*tag= a /mod_base= OTHER /note= "Biotin labelled"

Location/Qualifiers

modified base

Homo sapiens

Human telomerase RNA antisense oligonucleotide, oligo 5.

(first entry)

11-MAR-2004

ADG62872;

ADG62872 standard; DNA; 30 BP

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The invention describes a method of obtaining mammalian telomerase

protein (1). The method involves preparing enriched solution (ES) from a cell expressing telomerase where the component of (1) in ES is separated from other proteins expressed by cell by combining ES with coligonucleotide (0) having specific affinity for (1), and collecting protein bound to (0). The oligonucleotide comprises a retrievable label cut as biotin and contains a sequence that is specifically recognized by telomerase protein. The oligonucleotide comprises combining a fraction containing telomerase protein with an anion exchange matrix, and collecting protein that binds the matrix, combining a fraction containing telomerase protein that binds the matrix. The method collecting protein that binds the matrix. The method containing telomerase protein with a cation exchange matrix. The method containing telomerase protein that binds the matrix, where the intermediate containing telomerase protein with an intermediate selectivity matrix, collecting protein with an intermediate selectivity matrix, collecting protein with an intermediate selectivity matrix, collecting protein with an intermediate selectivity matrix, especially protein with an intermediate selectivity matrix, and separating a fraction containing the telomerase protein by gel filtration chromatography or gradient centrifugation. The telomerase is enriched from an extract of cells stably expressing telomerase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Obtaining telomerase, by preparing enriched solution from cell expressing telomerase, combining the solution with oligonucleotide having specific affinity for the protein and collecting protein bound to oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              telomerase. This sequence represents an antisense oligonucleotide to the RNA component of human telomerase that can be used in the purification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Weinrich SL, Atkinson EM, Lichtsteiner SP, Vasserot AP, Pruzan RA;
                                                                                                                                                      mammalian telomerase protein; telomerase purification; telomere; anton exchange matrix; selectivity matrix; gal filtration chromatography; gradient centrifugation; antisense oligonucleotide; ss.
                                                                                                            Human telomerase RNA component antisense oligonucleotide seq id
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 30 BP; 4 A; 11 C; 9 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 4; SEQ ID NO 3; 24pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                   95US-00510736.
97US-00833377.
99US-00420056.
                                                                                                                                                                                                                                                                                                                                                                                                           20-NOV-2000; 2000US-00717829
                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             method of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (GERO-) GERON CORP.
                                                                 18-DEC-2003
                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                 US6545133-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                       04-AUG-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             04-APR-1997;
18-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                            08-APR-2003
                       ADC35650;
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Pruzan RA;

Vasserot AP,

Lichtsteiner SP,

Weinrich SL, Atkinson EM,

WPI; 2003-811733/76.

WEINRICH S L.
ATKINSON E M.
LICHTSTEINER S P
VASSEROT A P.

(LICH/) (VASS/) (PRUZ/)

(WEIN/)

04-APR-1997; 97US-00833377 18-OCT-1999; 99US-00420056 20-NOV-2000; 2000US-00717828

04-AUG-1995; 04-APR-1997;

95US-00510736

24-DEC-2002; 2002US-00330872

US2003186282-A1

02-OCT-2003

Identifying telomerase regulators useful for treating cancer

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to a method for identifying regulators of telomerase activity that may be useful for treating cancers. The method may be used to identify regulators e.g. antibodies, of telomerase activity which may be useful as cancer treatments. It has been found that found that the cells of many human cancers have telomerase activity. This helps explain why cancer cells continue dividing without becoming senescent. If telomerase activity in cancer cells can be inhibited, the cancer cells are expected to reach senescence and cease dividing. The present sequence is human telomerase antisense oligonulceotide used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gарв
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ô
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       6.3%; Score 28.4; DB 1; Length 30; 96.7%; Pred. No. 45; lve 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 30 BP; 4 A; 11 C; 9 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         412 GAGCTGTGGGACGTGCACCCAGGACTCGGC 441
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GAGCTATGGGACGTGCACCCAGGACTCGGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              illustrate the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 11; SEQ ID NO 3; 22pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ilarity 96.7%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAX90787 standard; DNA; 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        13-JAN-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
tes 29; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAX90787;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Loc
Matches
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AC AAX9
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DT 13-J
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Gaps

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412 GAGCTGTGGGACGTGCACCCAGGACTCGGC 441

29; Conservative

Query Match Best Local Similarity Matches 29; Conserv

GAGCTATGGGACGTGCACCCAGGACTCGGC 1

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ADG62872/c

DB 1; Length 30; 1; Indels

6.3%; Score 28.4; D 96.7%; Pred. No. 45; iive 0; Mismatches

US09436060A-16.rng.sl

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This primer is used for the RT-PCR amplification of an RNA component of human telomerase (hTR). This is used in the method of invention of determining the amount of hTR in a sample. The method comprises camplifying a sequence of hTR and a control polynucleotide from a sample and determining an amount of amplified hTR and an amount of amplified control polynucleotide. The amount of amplified hTR is normalised with crespect to the amount of hTR which provides antisense oligonucleotides to creapect to the amount of hTR which provides antisense oligonucleotides to the hTR which may specifically be used for detection of an RNA component of hTR which may specifically be used for detection of an RNA component of human telomerase in a sample. This is useful for diagnosing cancer (especially neuroblastoma, bladder, colon and prostate cancer), and providing prognosis for a cancer patient. The antisense oligonucleotides con be used for inhibiting telomerase activity in both cultured cells and control of the immunosuppression, and for selectively down-regulating specific branches of the immune system, e.g. a specific subset of T-cells, in anti-nilammatory theraples or for treating infections by, e.g. yeast,
                                                                                                                                 New polynucleotide(s) anti:sense to human telomerase - used for detecting or inhibiting human telomerase, e.g. for treating cancers, contraception, immuno-suppression or treating infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New human GC6 gene, useful for identifying agents for treating diseases and/or conditions associated with cell senescence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cellular senescence; modulator; GC6 gene; senescent gene expression;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ·.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   6.0%; Score 27; DB 1; Length 27;
.00.0%; Pred. No. 51;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Indels
                Pruzan R, Weinrich SL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 27 BP; 7 A; 3 C; 10 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Pred. ...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 144 CCTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      27 ccrrccaccerrcarrcradaccaaac 1
                                                                                                                                                                                                                                                                     Claim 65; Page 75; 80pp; English
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                Kim NW, Wu F, Kealey JT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAX77130 standard; DNA; 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
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Matches 27; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PCR primer hTR445 comp
                                                                            WPI; 1998-377670/32.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               03-AUG-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAX77130;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAX77130/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present sequence is a PCR primer specific to human telomerase RNA (hTR). It is used to amplify human staufen (hStau) cDNA synthesised using random hexamer primers and Superscript II reverse transcriptase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RNA component; human telomerase; antisense oligonucleotide; infection; neuroblastoma; bladder cancer; colon cancer; prostate cancer; cancer; contraception; sterilisation; immunosuppression; therapeutic; hTR; immune system down-regulation; anti-inflammatory therapy; RT-PCR; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human staufen polypeptide useful in methods for identifying telomerase inhibitors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                  PCR primer; human telomerase RNA; hTR; amplify; human staufen cDNA; hStau; synthesised; random hexamer primer; Superscript II reverse transcriptase; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RNA component of human telomerase (hTR) amplifying reverse primer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 28 BP; 2 A; 4 C; 13 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (UYJO ) UNIV JOHNS HOPKINS SCHOOL MEDICINE.
Human telomerase RNA specific PCR primer-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17 GCCTGGGAGGGTGGTGGCCATTTTTG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAV41193 standard; DNA; 27 BP
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96US-00770565
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                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                                                                            WO9951255-A1
                                                                                                                                                                                                                                                                                                                                                                                                                    06-APR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Greider CW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          08-OCT-1998
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Matches

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Length 27;

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                                                                 senescence. Recombinant expression vectors comprising a recombinant polynucleotide corresponding to a polynucleotide in a human GG gene, are useful for altering senescent gene expression. The vectors and host cells comprising the vectors are useful for identifying agents that prevent or modulate senescent gene expression. The polynucleotides are useful for producing the protein, pGGG and nucleic acid derivatives. The proteins encoded are useful for raising antibodies specific for pGGG, which are useful for isolating pGGG, and for detecting cells comprising pGGG in complex cell mixtures. The characterization of the polynucleotides enable the identification of therapeutic agents that identify and distinguish between young and senescent cells. This enables treatment of aging diseases induced or exacerbated by cellular senescence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to a method for diagnosing and/or monitoring the evolution of cancers. The method comprises analysing enzyme telomerase RNA in blood plasma or serum. The method is typically used for diagnosing breast, ovarian, stomach or colon cancer and/or monitoring the evolution of the cancers after treatment by chemotherapy or operations. The present sequence is a PCR primer for human telomerase RNA (HTR), which was used in the example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     e.g. breast
plasma or serum.
                                                    The invention relates to methods for modulating and identifying cellular
                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; telomerase RNA; PCR primer; cancer; breast; ovarian; stomach;
colon; hTR; ss.
                                                                                                                                                                                                                                                                                                                                                                                ;
0
                                                                                                                                                                                                                                                                                                                                       Query Match 6.0%; Score 27; DB 1; Length 27; Best Local Similarity 100.0%; Pred. No. 51; Matches 27; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         e, reliable diagnosis and/or prognosis of cancer, by analyzing the RNA components of telomerase in
                                                                                                                                                                                                                                                                                                         Sequence 27 BP; 4 A; 6 C; 11 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human telomerase RNA, hTR, antisense PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                  425 TGCACCCAGGACTCGGCTCACACATGC 451
                                                                                                                                                                                                                                                                                                                                                                                                                                     27 TGCACCCAGGACTCGGCTCACACATGC 1
                  5; Page 74; 79pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABA95497 standard; DNA; 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-099090/14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (CHEN/) CHEN X Q. (STRO/) STROUN M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ANKER P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABA95497;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 49
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Sequence 27 BP; 7 A; 3 C; 10 G; 7 T; 0 U; 0 Other;

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The invention relates to promoter regions from mouse and human telomerase

RNA (TR) component genes. The TR gene promoter can be linked to a

RNA (TR) component genes. The TR gene encoding a cytocoxin, for therapy of

cancer, especially neoplasias. The telomerase is necessary for the

cancer, especially neoplasias. The telomerase is necessary for the

currestricted proliferative capacity of many human cancers. Mutation or

dysregulation of the telomerase repression pathway may cause reactivation

or upregulation of telomerase expression in cancer. Substances,

cleantified in the methods, can be used to block transcription from the TR

consideration of telomerase expression in cancer. Substances.

Clean promoter through interaction of the 5' regulatory sequences. These

substances, e.g. antisense oligonucleotides, transcription factors,

cuseful for cancer therapy. In particular, gene therapy vectors

cuseful for cancer therapy. In particular, gene therapy vectors

cuseful for cancer therapy. In particular, gene therapy vectors

cuseful for cancer therapy. In particular, gene therapy vectors

cuseful for cancer therapy. In particular, gene therapy vectors

cuseful for cancer therapy and the promoter and a viral thymidine

kinase gene can be controlled or treated. Direct down-regulation of

ceffective anticancer therapy and the cloning of the hTR gene promoter

allows the analysis of therapeutic molecules which modulate hTR promoter

callows the analysis of therapeutic molecules which modulate hTR promoter

callows the analysis of therapeutic molecules which modulate for amplifying
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mouse and human telomerase RNA gene promoters, useful for tumor specific
                                            Gape
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gape
                                                                                                                                                                                                                                                                                                                                                                                                          Telomerase RNA; TR; promoter; cytotoxin; cancer; neoplasia; hTR; gene therapy; thymidine kinase gene; anticancer therapy; human;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                5.9%; Score 26.8; DB 1; Length 30; 93.3%; Pred. No. 61; cive 0; Mismatches 2; Indel8
                                          0; Indels
                                                                                                                                                                                                                                                                                                                                                                Human telomerase RNA gene (hTR) specific primer hTR14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 30 BP; 5 A; 8 C; 8 G; 9 T; 0 U; 0 Other;
6.0%; Score 27; DB 1
ilarity 100.0%; Pred. No. 51;
Conservative 0; Mismatches
                                                                                    144 CCTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (CANC-) CANCER RES CAMPAIGN TECHNOLOGY.
                                                                                                                   CCTTCCACCGTTCATTCTAGAGCAAAC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  promoter sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Fig 6; 109pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           98GB-00001902
                                                                                                                                                                                                                    264/c
AAZ07264 Btandard; DNA; 30
                                                                                                                                                                                                                                                                                                                        22-OCT-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1999-479183/40
Query Match
Best Local Similarity
Matches 27; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity
Matches 28, Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                         PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                human TR gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           29-JAN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   29-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             W09938964-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       gene therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         05-AUG-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
                                                                                                                                                                                                                                                                                AAZ07264;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Keith WN;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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mammal; antisense; triplex forming oligonucleotide; plasmid;

(first entry)

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The RNA components of (non) human mammalian telomerase(s) especially from mouse, rat and chinese hamster are all claimed. Antisense oligonucleotides can be used to block the activity of the telomerase; probes and primers can be used in detection; vectors and host cells transformed with the isolated telomerase genes can be used for production of telomerases; RNA and DNA ribozymes and triplex forming of telomerases in the rate of the solated telomerase genes can be used the reported against the telomerase genes can be used the rapeutically as can plasmids. A mouse which lacks the telomerase gene (also claimed) can be used for study of telomera regulation in vivo, and the role it plays in immortalisation. Three primers (AAT11040, AAT11041).
                                                                Primer for production of telomerase antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   senescence and immortalisation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 8; Page 53; 85pp; English.
                                                                                                             probe; primer; ribozyme; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1996-097428/10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  RNA components of
                                                                                                                                                                                                                                                                                                                                                                                         Villeponteau B;
                                    02-JUL-1996
                                                                                                                                                                    WO9601614-A2
                                                                                                                                                                                                                            07-JUL-1995;
                                                                                                                                                                                                                                                         07-JUL-1994;
                                                                                                                                                                                                                                                                       27-OCT-1994;
13-FEB-1995;
                                                                                                                                                                                                                                                                                                    07-JUN-1995;
                                                                                             Telomerase;
                                                                                                                                                                                                25-JAN-1996
                                                                                                                                                                                                                                                                                                                                                                           Andrews WH,
                                                                                                                                       Synthetic.
         AAT11044;
                                                                                                                                                                                                                                                                                                                                (COLD-)
                                                                                                                                                                                                                                                                                                                                               GERO-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present sequence, a forward primer for the RNA component of human telomerase (RCHT), was used in a primed in situ labelling (PRINS) procedure. The RCHT can be used in the recombinant prodn. of an active telomerase mol., capable of adding sequences to chromosomal DNA telomeras, and in the synthesis of mutant sequences for the detection of mutant mammalian telomerase RNA component polymoleotides. The RCHT may also be used in the identification of telomerase modulating agents, and in the detection of telomerase related, or neoplastic conditions in a patient. Polymoleotides of at least 25 consecutive mucleotides identical, or complementary to the RCHT sequence linked to heterologous transcriptional regulatory sequences, can be used for the gene therapy of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RNA component of mammalian telomerase, esp. human - useful in identifying
                                                                                                                                                                                                                   RNA component; human; telomerase; forward primer; PRINS; recombinant production; synthesis; mutant; detection; mammalian; identification; modulating agent; neoplastic condition; transcriptional regulatory sequence; gene therapy; disease; primed in situ labelling; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1; Length 28;
                                                                                                                                                                                       RNA component of human telomerase PRINS forward primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 28 BP; 2 A; 3 C; 13 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Andrews WH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          e.g. candidate telomerase-modulating agents.
                  CTAACCCTAACTGAGAAGGGCGTAGGATCC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 26.4;
Pred. No. 60
CTAACCCTAACTGAGAAGGGCGTAGGCGCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 13; Page 91; 114pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Funk W,
                                                                                                     BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                         94US-00330123.
95US-00472802.
95US-00482115.
                                                                                                                                                                                                                                                                                                                                                                                                  95WO-US008530
                                                                                                                                                                                                                                                                                                                                                                                                                              94US-00272102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5.9%;
                                                                                                  AAT10309 standard; DNA; 28
                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Feng J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1996-097581/10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (GERO-) GERON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Villeponteau B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               human diseases
                                                                                                                                                                                                                                                                                                                                        WO9601835-A1
                                                                                                                                                                                                                                                                                                                                                                                                 06-JUL-1995;
                                                                                                                                                             10-SEP-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                            27-OCT-1994;
                                                                                                                                                                                                                                                                                                                                                                   25-JAN-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-JUN-1995
                                                                                                                                                                                                                                                                                                            Synthetic.
                          30
                                                                                                                               AAT10309;
 46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                     RESULT 51
                                                                                                                à
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(non)human mammalian telomerase(s) - useful in studying

Marhuenda MA;

Greider C,

Funk W,

Feng J,

Avillon AA,

94US-00272102. 94US-00330123. 95US-00387524. 95US-00485778.

95WO-US008620

SPRING HARBOR LAB

GERON CORP

COLD

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used to produce antisense expression plasmids. AAT11040 was used alongside both AAT11043 and AAT11044 to produce two different antisense
                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                   RNA component; human; telomerase; polymerase chain reaction; recombinant production; synthesis; mutant; detection; mammalian;
                                                                                                                                            ö
                                                                                                      Score 26; DB 1; Length 26; Pred. No. 59;
                                                                                                                                          0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                  RNA component of human telomerase nested PCR primer R3c.
                                                                     Sequence 26 BP; 7 A; 3 C; 9 G; 7 T; 0 U; 0 Other;
                                                                                           5.8%; Sco. 100.0%; Pred. No. ... 0; Mismatches
                                                                                                                                                                           145 CTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                            26 crrccaccerrcarrcragagcaaac 1
                                                                                                                                                                                                                                                                                                   BP
                                                                                                                                                                                                                                                                                 304/c
AAT10304 standard; DNA; 26
                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                        26; Conservative
                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                   10-SEP-1996
                                      molecules
                                                                                                                                                                                                                                                                                                                                  AAT10304;
                                                                                                       Query Match
                                                                                                                          Local
                                                                                                                                                                                                                                                                            AAT10304/c
ID AAT1C
XX
AC AAT1C
XX
XX
DT 10-SE
XX
CX
KW RNA C
KW RWA C
                                                                                                                                                                                                                                                                 RESULT 53
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0

Gaps

. 0

1; Indels

0; Mismatches

Conservative

Local Similarity les 27; Conser

Matches

44

GCTGGGAGGGTGGTGGCTATTTTTG 17 GCCTGGGAGGGGTGGTGGCCATTTTTG

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AAT11044 standard; DNA; 26 BP

AAT11044/c ID AAT110 XX

RESULT 52

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(GERO-) GERON
                                      07-JUL-1994;
27-OCT-1994;
07-JUN-1995;
07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO9601835-A1.
  06-JUL-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           06-JUL-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    07-JUL-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         27-OCT-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  07-7UN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25-JAN-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAT10306;
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present sequence, a nested PCR primer for the RNA component of human telomerase (RCHT), was used in a 5' RACE procedure. The RCHT can be used in the recombinant prodn. of an active telomerase mol., capable of adding sequences to chromosomal DNA telomeras, and in the synthesis of mutant sequences for the detection of mutant mammalian telomerase RNA component polynucleotides. The RCHT may also be used in the identification of telomerase modulating agents, and in the detection of telomerase related, or neoplastic conditions in a patient. Polynucleotides of at least 25 consecutive nucleotides identical, or complementary to the RCHT sequence linked to heterologous transcriptional regulatory sequences, can be used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RNA component of mammalian telomerase, esp. human - useful in identifying
modulating agent; neoplastic condition; regulatory sequence; gene therapy; disease; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RNA component; human; telomerase; lung fibroblast; cell line Wi 38; recombinant production; synthesis; mutant; detection; mammalian; dentification; madulating agent; neoplastic condition; transcriptional regulatory sequence; gene therapy; disease; polymerase chain reaction; antisense plasmid; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RNA component of human telomerase antisense plasmid PCR primer R3C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5.8%; Score 26; DB 1; Length 26;
100.0%; Pred. No. 59;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                           Funk W, Andrews WH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               e.g. candidate telomerase-modulating agents.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Preα. ...
.... 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      145 CTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    gene therapy of human diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 10; Page 82; 114pp; English
                                                                                                                                                                                                                                                               94US-00330123.
95US-00472802.
95US-00482115.
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                                                                                                                                                                                                                                                                                                                                                                                      Villeponteau B, Feng J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1996-097581/10.
                                                                                                                                                                                                                                                                                                                                                  (GERO-) GERON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
hes 26; Conserv
identification;
transcriptional
                                                                                                                       WO9601835-A1
                                                                                                                                                                                                      06-JUL-1995;
                                                                                                                                                                                                                                                                                                          07-JUN-1995;
                                                                                                                                                                                                                                              07-JUL-1994;
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07-JUN-1995;
                                                                                                                                                               25-JAN-1996
                                                                                Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            56
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KEWERE ENGLAND TO SEE THE SEE 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ઠે
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Polynucleotides of at least 25 consecutive nucleotides identical, or
                                                                                                                                                                                                                                                                                                                                                                                                                         RNA component of mammalian telomerase, esp. human - useful in identifying
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                complementary to the RCHT sequence linked to heterologous transcriptional regulatory sequences, can be used for the gene therapy of human diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RNA component; human; telomerase; polymerase chain reaction; recombinant production; synthesis; mutant; detection; mammalian; dentification; modulating agent; neoplastic condition; transcriptional regulatory sequence; gene therapy; disease; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gapв
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        5.8%; Score 26; DB 1; Length 26;
100.0%; Pred. No. 59;
iive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 26 BP; 7 A; 3 C; 9 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RNA component of human telomerase PCR primer F3b.
                                                                                                                                                                                                                                                                                         Andrews WH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                           e.g. candidate telomerase-modulating agents.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                145 CTTCCACCGTTCATTCTAGAGCAAC 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CTTCCACCGTTCATTCTAGAGCAAAC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 8; Page 80; 114pp; English.
                                                                                                                                                                                                                                                                                         Funk W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  94US-00272102.
94US-00330123.
95US-00472802.
95US-00482115.
                                                         94US-00272102.
94US-00330123.
95US-00472802.
95US-00482115.
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95WO-US008530.
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Best Local Similarity 100.
                                                                                                                                                                                                                                                                                         Feng J,
                                                                                                                                                                                                                                                                                                                                                            WPI; 1996-097581/10.
                                                                                                                                                                                                                            CORP.
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                                                                                                                                                       The present sequence, a PCR primer for the RNA component of human telomerase (RCHT), was used in a 3' RACE procedure. The RCHT can be used in the recombinant prodn. of an active telomerase mol., capable of adding sequences to chromosomal DNA telomerase, and in the synthesis of mutant sequences for the detection of mutant mammalian telomerase RNA component polynucleotides. The RCHT may also be used in the identification of telomerase modulating agents, and in the detection of telomerase related, or neoplastic conditions in a patient. Polynucleotides of at least 25 consecutive nucleotides identical, or complementary to the RCHT sequence linked to heterologous transcriptional regulatory sequences, can be used
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                                                                  mammalian telomerase, esp. human - useful in identifying
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cancer; eukaryotic parasite; hTR; vertebrate telomerase; yeast; protozoa; tumour; antibody; polymerase chain reaction; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA encoding essential RNA components of human telomerase - also truncated or recombinant telomerase, useful for diagnosis and treatment of cancer and infection by eukaryotic parasites.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present sequence represents PCR 3'-primer R3C used for amplifying human telomerase (hTR). The RNA and DNA can be used in hybridisation assays to detect or quantify telomerase activity in cells, tissue or fluid samples, e.g. for diagnosis of eukaryotic parasites (yeast and protozoa) or tumours. It is also useful as primers for amplification assays. The truncated or recombinant vertebrate telomerase is used
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100.0%; Pred. No. 59;
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                                                                                                                                                                                                                                                                                                                                                                          Seguence 26 BP; 8 A; 6 C; 7 G; 5 T; 0 U; 0 Other;
 Andrews WH;
                                                                    RNA component of mammalian telomerase, esp. e.g. candidate telomerase-modulating agents.
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                                                                                                                                                                                                                                                                                                                                       for the gene therapy of human diseases
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                                                                                                                          Example 10; Page 83; 114pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human telomerase PCR 3'-primer R3C.
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Funk W,
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 Villeponteau B, Feng J,
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                                   WPI; 1996-097581/10
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therapeutically to increase telomerase activity (also as reagents in the screening assay) while the RNA or other inhibitors such as antisense molecules, are used to reduce such activity. Typical applications are initiation/restoration of activity to cause senescence or to prevent immortalisation of cells in tumours or parasites. The DNA is also used to produce recombinant telomerase, which can then be used conventionally to raise antibodies for diagnostic detection of telomerase. Detecting telomerase allows early diagnostic detection of telomerase. Detecting signs manifest. Telomerase inhibitors directed against e.g. Trypanosoma should cause fewer side effects than drugs currently used to treat such infections. The DNA encodes those parts of hTR RNA essential for activity but are significantly shorter than the endogenous RNA component
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RNA component; human telomerase; antisense oligonucleotide; infection; neuroblastoma; bladder cancer; colon cancer; prostate cancer; cancer; contraception; sterilisation; immunosuppression; therapeutic; hTR; immune system down-regulation; anti-inflammatory therapy; RT-PCR; primer;
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100.0%; Pred. No. 59;
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                                                                                                                                                                                                                                                                                                                                                                                             145 CTTCCACGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                                                                                                                                                                                        26 CTTCCACCGTTCATTCTAGAGCAAAC 1
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nes 26; Conservative
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AAV41192
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the hTR which may specifically be used for detection of an RNA component of human telomerase in a sample. This is useful for diagnosing cancer (especially neuroblastoma, bladder, colon and prostate cancer), and providing prognosis for a cancer patient. The antisense oligonucleotides can be used for inhibiting telomerase activity in both cultured cells and in cells in vivo. They can be used in therapeutics for treating or immunosuppression, and for selectively down-regulating specific branches of the immune system, e.g. a specific subset of T-cells, in anti-inflammatory therapies or for treating infections by, e.g. yeast,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human, telomerase reverse transcriptase, hTRT, TRT, diagnosis, prognosis, cell proliferation, cancer, ageing, ribonucleoprotein, PCR primer, ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present sequence represents a PCR primer from the present invention which describes human telomerase reverse transcriptase (hTRT). The present invention also describes the following methods. (A) determining whether a test compound is a modulator of hTRT, by detecting the change in hTRT recombinant protein or polynucleotide, on administration of the
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                                                                                                                                                                                                                                                                         60 GAAGGCCTAGGCGCCGTGCTTTTGC
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970S-00846017.
970S-00851843.
970S-0091312.
970S-0091312.
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ID AAV17033 standard; DNA; 26
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compound; (B) preparation of recombinant telomerase by contacting a protein preparation of HTRT with a telomerase RNA component; (C) detection of the HTRT RNA or protein in a sample by binding a relevant probe to the sample and detecting the complex formed or in the case of RNA detection, amplifying the product and correlating the presence of complex or amplification product with presence of hTRT in the sample, and (D) increasing the proliferation of a vertebrate cell by increasing hTRT expression; and (E) the use of an agent that causes an increase in cell vertebrate cell proliferation to create a medicament that inhibits ageing. A protein preparation of hTRT and the polynucleotide encoding hTRT can be used in the manufacture of medicaments for inhibiting the effect of ageing or cancer. Inhibitors of telomerase activity can be used to treat conditions that are associated with high telomerase activity. A protein preparation of hTRT can also be used in the new methods
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100.0%; Pred. No. 59;
tive 0; Mismatches
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97US-00846017.
97US-00851843.
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UNIV TECHNOLOGY CORP.
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Best Local Similarity 100.
Matches 26; Conservative
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present invention also describes the following methods: (A) determining whether a test compound is a modulator of hTRT, by detecting the change in hTRT recombinant protein or polymucleotide, on administration of the compound; (B) preparation of recombinant telomerase by contacting a protein preparation of the hTRT with a telomerase BNA component; (C) detection of the hTRT RNA or protein in a sample by binding a relevant probe to the sample and detecting the complex formed or in the case of RNA detection, amplifying the product and correlating the presence of complex or amplification product with presence of hTRT in the sample; and (D) increasing the proliferation of a vertebrate cell by increasing hTRT expression; and (E) the use of an agent that causes an increase in cell vertebrate cell proliferation of reate a medicament that inhibits of ageing A protein preparation of hTRT and the polymucleotide encoding hTRT can be used in the manufacture of medicaments for inhibiting the effect of ageing or cancer. Inhibitors of telomerase activity can be used
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                                                                                                                                                                                                                                                                                     treat conditions that are associated with high telomerase activity. A
                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human TPC2, TPC3 and TR genes - regulate telomere length or modulate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              gene; TPC2; TPC3; telomere length; telomerase; human; cancer;
                                                                                                                                                                                                                                                                                                    protein preparation of hTRT can also be used in the new methods
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 49; 86pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human hTR gene RT-PCR primer F3b.
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Best Local Similarity 100.
Best Aca 26; Conservative
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                             Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               TPC3 and TR genes - regulate telomere length or modulate
                                                                                                                                                                                                                            hTR gene; TPC2; TPC3; telomere length; telomerase; human; cancer; gene therapy; diagnosis; PCR; primer; ss.
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59;
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      DB 1;
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                                                    45 TCTAACCCTAACTGAGAAGGGCGTAG
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AAX90788 standard; DNA; 26
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       Query Match 5.8
Best Local Similarity 100.
Matches 26; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                              (GERO-) GERON CORP.
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                                                                                                                                                          AAV19489;
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telomeric repeat amplification protocol.
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                                                                                                                                                                                                                            The present sequence is a PCR primer specific to human telomerase RNA (hTR). It is used to amplify human staufen (hStau) cDNA synthesised using random hexamer primers and Superscript II reverse transcriptase
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                                                                                                                                                                                    Human staufen polypeptide useful in methods for identifying telomerase
                                                                                                                                                                                                                                                                                               Gaps
     PCR primer; human telomerase RNA; hTR; amplify; human staufen cDNA; hStau; synthesised; random hexamer primer; Superscript II reverse transcriptase; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                            Telomerase; body fluid; cancer; tumour; screening; TRAP; diagnosis; telomeric repeat amplification protocol; detection; PCR primer; ss.
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                                                                                                                                                                                                                                                              Sequence 26 BP; 7 A; 3 C; 9 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                             Human telomerase RNA template PCR primer F3B.
                                                                                                                                  (UYJO ) UNIV JOHNS HOPKINS SCHOOL MEDICINE
                                                                                                                                                                                                                                                                                                                 145 CTTCCACCGTTCATTCTAGAGCAAAC 170
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                                                                                                                                                                                                                                                                                                                         CTTCCACCGTTCATTCTAGAGCAAAC 1
                                                                                                                                                                                                             Disclosure; Page 15; 50pp; English
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Best Local Similarity
Matches 26; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                            Telomerase; body
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                                                 Homo sapiens
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                                                                                                  36-APR-1999;
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                                                                                                                                                   Greider CW,
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                                                                                                                                                                                            inhibitors
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                                        Synthetic
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cellular portion of body fluid from a cancer patient using a modified cellular portion of body fluid from a cancer patient using a modified tellular portion of body fluid from a cancer patient using a modified cellular portion of a modified cancer comprises: (a) removing the cellular portion of a body fluid specimen from the patient; (b) preparing a protein extract from the body fluid remainder; (c) assaying the extract for the presence and quantity of telomerase RNA or telomerase activity; and (d) comparing the results with normal levels, to determine the presence of cancer. The methods are therapy effectiveness Unlike prior art telomerase activity assays in cancer patients, the method allows noninvasive sample collection. The methods are also more reliable and less tumour specific than other methods which detect circulating tumour markers. The present sequence represents a human telomerase RNA template PCR primer used in the
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Disclosure; Page 16; 32pp; English
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Best Local Similarity
Matches 26; Conserv
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pGC6; human; PCR primer; ss.
                                                                                               (GERO-) GERON CORP
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                                                                                19-NOV-1997;
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                                                                19-NOV-1998;
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                                                27-MAY-1999
                 Synthetic.
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                         Gaps
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                                                                                                                                                             Telomerase; human; diagnosis; bladder cancer; detection; urine;
PCR primer; ss.
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        DB 1; Length 26;
59;
                         0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 26 BP; 7 A; 3 C; 9 G; 7 T; 0 U; 0 Other;
             100.0%; Prec. ...
                                                                                                                                                  Human telomerase RNA PCR primer TE-hTR3.1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           145 CTTCCACCGTTCATTCTAGAGCAAAC 170
          5.8%; Score 26;
100.0%; Pred. No.
                                         79
                                                    1 AACTGAGAAGGGCGTAGGCGCCGTGC 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CTTCCACCGTTCATTCTAGAGCAAAC 1
                                          54 AACTGAGAAGGGCGTAGGCGCGTGC
                                                                                                                                                                                                                                                                                                                                                           Claim 6; Page 10; 13pp; German.
                                                                                                                                                                                                                                                                                   (HOFF ) ROCHE DIAGNOSTICS GMBH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAX77131 standard; DNA; 26 BP
                                                                                                  BP
                                                                                                                                                                                                                                                                   97DE-01057300.
                                                                                                                                                                                                                                                   98EP-00124326
                                                                                                  AAX77402 standard; DNA; 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                  (first entry)
                          Conservative
                                                                                                                                                                                                                                                                                                                   WPI; 1999-349242/30.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PCR primer hTR S328
                  Best Local Similarity
                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                   21-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    03-AUG-1999
                                                                                                                                                                                                                                                                   22-DEC-1997;
                                                                                                                                  05-AUG-1999
                                                                                                                                                                                                                    EP926245-A2.
                          26;
                                                                                                                                                                                                                                   30-JUN-1999
                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                                   Emrich T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAX77131;
                                                                                                                  AAX77402;
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          Query Match
                          Matches
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The invention relates to methods for modulating and identifying cellular senescence. Recombinant expression vectors comprising a recombinant polymucleotide corresponding to a polymucleotide in a human GGG gene, are useful for altering senescent gene expression. The vectors and host cells comprising the protein, pGGG and mucleic acid derivatives. The prevent or modulate senescent gene expression. The polymucleotides are useful for porducing the protein, pGGG and mucleic acid derivatives. The proteins encoded are useful for raising antibodies specific for pGGG, which are useful for isolating pGGG, and for detecting cells comprising pGGG in complex cell mixtures. The characterization of the polymucleotides enable the identification of therapeutic agents that identify and distinguish between young and senescent cells. This enables treatment of aging diseases induced or exacerbated by cellular senescence
                                                                                                                                                                                                                                                                                                                                New human GCG gene, useful for identifying agents for treating diseases and/or conditions associated with cell senescence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TPC2, TPC3; human; telomere length regulation; cancer; pregnancy; fertility; diagnosis; therapy; PCR primer; ss.
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59;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Pred. w.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  306 TIGGGCTCTGTCAGCCGCGGGTCTCT 331
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                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 5; Page 74; 79pp; English
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96US-00583808.
98WO-US024996.
                                                                97US-00974180.
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Best Local Similarity 100.
Matches 26; Conservative
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WO200046601-A1
                                                                                                                                                                                                                                                                                                                                         32-FEB-1999;
                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                   15-DEC-2000
         Sequence 26
                                                                                                                                                                                                                                                                                                   10-AUG-2000
                                                                                         56
                                                                                                                                                                AAA88250;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                             Human;
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                                                                                                                       RESULT 69
                                                                                                                                   AAA88250,
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                                                                                                           This sequence represents a PCR primer for DNA encoding the human TPC3 protein, which is contained within the recombinant mammalian host cell of the invention. The invention provides methods and reagents for regulating telomerae and reagents for regulating telomerae activity in mammalian cells as well as for detecting, diagnosing, and treating related diseases and conditions such as cancer, pregnancy, or fertility in humans and other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This sequence represents a PCR primer for DNA encoding the human TPC3 protein, which is contained within the recombinant mammalian host cell of the invention. The invention provides methods and reagents for regulating telomerae activity in mammalian cells as well as for detecting diagnosing, and treating related diseases and conditions such as cancer, pregnancy, or fertility in humans and other
                                                                                                                                                                                                                                      Gaps
                                                           DNA encoding proteins TPC2 and TPC3 - useful for regulating telomere length or modulating telomerase activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA encoding proteins TPC2 and TPC3 - useful for regulating telomere length or modulating telomerase activity.
                                                                                                                                                                                                                                                                                                                                                                                                               TPC2, TPC3; human; telomere length regulation; cancer; pregnancy; fertility; diagnosis; therapy; PCR primer; ss.
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0
                                                                                                                                                                                                                5.8%; Score 26; DB 1; Length 26;
100.0%; Pred. No. 59;
iive 0; Mismatches 0; Indels
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                    Feng
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                   Villeponteau B,
                                                                                                                                                                                                                                                           TCTAACCCTAACTGAGAAGGGCGTAG 70
                                                                                                                                                                                                                                                                            TCTAACCCTAACTGAGAGGGCGTAG 26
                                                                                         Example, Col 38; 59pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example; Col 38; 59pp; English.
                                                                                                                                                                                                                                                                                                                                                                                             PCR primer for Human TPC3 gene
                                                                                                                                                                                                                                                                                                                                 BP
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ID AAX01542 standard; DNA; 26
                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Andrews WH,
                    Andrews WH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1999-152104/13.
                                       WPI; 1999-152104/13
(GERO-) GERON CORP.
                                                                                                                                                                                                                            Local Similarity
nes 26; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (GERO-) GERON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                              TPC3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      13-SEP-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         08-SEP-1995;
05-JAN-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic
                    Adams RR,
                                                                                                                                                                                                                                                           45
                                                                                                                                                                                                                                                                                                                                                    AAX01542;
                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     nammals
                                                                                                                                                                                                                                                                                                             RESULT 68
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activity in a sample. The method comprises treating the sample with a solid phase to bind telomerase, separating the solid phase from the solid phase to bind telomerase, separating the solid phase from the sample which may be treated to elute bound telomerase and assaying the sample for telomerase activity. Also described are: (1) a kit (11) for detecting telomerase activity, and (2) a component (111) of an assay system for telomerase activity, and (2) a component (111) of an assay system for telomerase activity, comprising a solid phase for binding telomerase activity, comprising a solid phase for binding telomerase on which is present a substrate for telomerase elongation. (1) is useful for ancer diagnosis or prognosis and detection of micrometastasis. The solid phase used in (1) is useful for separating telomerase from a sample and therefore for detecting telomerase from a sample and therefore for detecting to telomerase activity. (11) is useful for detection of cancer calls and may also comprises means for assaying a mannal for assaying a mannal for assaying a reverse transcriptase (RT) PCR primer for
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Detecting telomerase activity in samples, useful for diagnosis of cancer and micrometastasis, comprises treating sample with solid phase, removing solid phase and treating to elute bound telomerase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   present invention describes a method (I) for detecting telomerase
                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               hTR; reverse transcriptase; RT-PCR; PCR primer; micrometastasis; diagnosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          telomerase RNA, which is used in an example from the present
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0
                                                DB 1; Length 26; 59;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human telomerase RNA reverse transcriptase PCR primer #2.
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100.0%; Pred. No. 59;
1ve 0; Mismatches 0; Indels
                                                                                                                        Indele
G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 26 BP; 7 A; 3 C; 9 G; 7 T; 0 U; 0 Other;
                                                                                 100.0%; Prec. ...
                                                             Score 26;
Pred. No.
                                                                                                                                                                                     145 CTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                   CTTCCACCGTTCATTCTAGAGCAAAC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 11; Page 38; 68pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                           BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%;
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   BP; 7 A; 3 C; 9
                                                             5.8%;
                                                                                                                                                                                                                                                                                                                                                                                                        AAA88250 standard; DNA; 26
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                                                                                                                           Conservative
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                                                   Query Match
Best Local Similarity
Matches 26; Conser
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les 26, Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  telomerase;
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(SKAA/) SKAANSENG
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The present invention describes a method (I) for detecting telomerase activity in a sample. The method comprises treating the sample with a solid phase to bind telomerase, separating the solid phase from the solid phase to bind telomerase, separating the solid phase from the telomerase and assaying the sample for telomerase activity. Also described are: (1) a kit (II) for detecting telomerase activity, comprising a solid phase and one or more components for assaying telomerase activity, comprising a solid phase activity, comprising a solid phase for binding telomerase activity of prognosis and detection of telomerase elongation. (I) is useful for ancer or micrometastasis. The solid phase used in (I) is useful for separating telomerase from a sample and therefore for detecting telomerase from a sample and therefore for detecting telomerase from a sample and therefore for cancer. The sparating telomerase for assaying an mRNA diagnostic for cancer. The present sequence represents a reverse transcriptase (RT) PCR primer for human telomerase RNA, which is used in an example from the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Detecting telomerase activity in samples, useful for diagnosis of cancer and micrometastasis, comprises treating sample with solid phase, removing solid phase and treating to elute bound telomerase.
                                                                                                                                                                                                                                                                                               telomerase; hTR; reverse transcriptase; RT-PCR; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5.8%; Score 26; DB 1; Length 26; 00.0%; Pred. No. 59;
                                                                                                                                                                                                                                                            Human telomerase RNA reverse transcriptase PCR primer #1.
                                                                                                                                                                                                                                                                                                     Human; telomerase; hTR; reverse transcriptase; RT-
detection; cancer; micrometastasis; diagnosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 26 BP; 8 A; 6 C; 7 G; 5 T; 0 U; 0 Other;
CTTCCACCGTTCATTCTAGAGCAAAC 170
                                   criccaccerrcarrcradadcaaac 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 11; Page 38; 68pp; English.
                                                                                                                                         ВР
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-FEB-2000; 2000WO-IB000100.
                                                                                                                                       AAA88249 standard; DNA; 26
                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Skaanseng M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (LARS/) LARSEN F.
(SKAA/) SKAANSENG M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2000-491281/43.
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                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
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                                                                                                                                                                                                                      15-DEC-2000
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                                                                                                                                                                                 AAA88249;
    145
                                         56
                                                                                                    RESULT 70
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The invention relates to detecting human telomerase RNA (FUERT) RNA (CC in a bodily fluid, comprising amplifying RNA extracted from plasma or serum sample, or its corresponding CDNA comprising (I), using primers or probes that target (I) or CDNA and detecting qualitatively or probes that target (I) or CDNA and detecting qualitatively or constitutively amplified product of (I) or CDNA product. The method is useful for detecting (I) in a bodily fluid, which is useful for constitutively amplified broader (I) expressing cells or tissue which include a malignancy preferably a cancer of breat, ovarian, head and neck, lung, cervical, colorectal, gastric, liver, pancreatic, bladder, prostate, condometrial, brain, kidney, or oesophagus, or amelanoma or sarcoma, cervical intrappthalial neophasia, bronchial dysplasia, atypical carcinoma in-situ, preferably cervical dysplasia, cartypical endometrial hyperplasia, or Barrett's oesophagus, where the hyperplasia of the breast, ductal carcinoma in-situ, colorectal adenoma, atypical endometrial hyperplasia, or Barrett's oesophagus, where the construction of the construction of the malignancy or premalignancy and for monitoring a therapy administered to a human. (I) provides a marker which is utilised as a during or following surgical procedures to remove the humans during or following surgical procedures to remove consulyment or malignant or menaphancy or qualitatively, of (I) in plasma or premalignant or menaphancy or relapsed cancer, impending relapse, or treatment failure.
                                                                                                                                                                                                                           cancer; breast; ovarian; head and neck; lung; cervical; colorectal; cancer; breast; ovarian; head and neck; lung; cervical; cosophagus; melanoma; sarcoma; premalignancy; carcinoma in-situ; cervical dysplasia; bronchial dysplasia; cervical intraepithelial neoplasia; atypical hyperplasia; colorectal adenoma; atypical endometrial hyperplasia; tumour; Barrett's oesophagus; telomerase-directed therapy; primer; ss.
                                                                                                                                                                                                     telomerase-associated RNA template; hTR; endometrial; malignancy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   reverse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to detecting human telomerase RNA template (hTR)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Detecting human telomerase RNA template RNA or human telomerase transcriptase protein RNA in bodily fluid, useful as marker for diagnosing, monitoring or treating cancer, carcinoma in situ or premalignancy.
                                                                                                                                                        Human telomerase-associated RNA template (hTR), PCR primer hTR2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 1; Page 14; 30pp; English.
                 ABK48024 standard; DNA; 26 BP.
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                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Kopreski MS, Gocke CD;
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ABK48024/c
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Gaps

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0; Indels

100.0%; Pred. No. 59; ive 0; Mismatches

Best Local Similarity 100. Matches 26; Conservative

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TCTAACCCTAACTGAGAAGGGCGTAG

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rcraacccraacrgagaagggggrag 26

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cancer; breast; ovarian; head and neck; lung; cervical; colorectal; gastric; liver; pancreatic; bladder; prostate; brain; kidney; oesophagus; melanoma; sarcoma; premalignancy; carcinoma in-aitu; cervical dysplasia; bronchial dysplasia; cervical intraspithelial neoplasia; atypical hyperplasia; colorectal adenoma; atypical endometrial hyperplasia; tumour; Barrett's oesophagus; telomerase-directed therapy; primer; ss.
The method allows the development and application of telomerase-specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; telomerase-associated RNA template; hTR; endometrial; malignancy;
                                     therapy even when only premalignant tumours, early cancer, or occult cancer or metaertasis auch as following resection or in minimal residual disease are present. The present sequence represents a PCR primer for human telomerase-associated RNA template (hTR)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           reverse
                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human telomerase-associated RNA template (hTR), PCR primer hTR1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Detecting human telomerase RNA template RNA or human telomerase transcriptase protein RNA in bodily fluid, useful as marker for diagnosing, monitoring or treating cancer, carcinoma in situ or
                                                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                                                                                                                                                                                                    5.8%; Score 26; DB 1; Length 26; 100.0%; Pred. No. 59;
                                                                                                                                                                                                                                                                                                                                                                   0; Indels
                                                                                                                                                                                                                    Sequence 26 BP; 7 A; 3 C; 9 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                    145 CTTCCACGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26 criccaccericaricradagedade 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABK48023 standard; DNA; 26 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       31-AUG-2000; 2000US-00653573.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28-AUG-2001; 2001WO-US026749
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                                                                                                                                                                                                                                                                                                                      Local Similaricy
nes 26; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2002-269532/31.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO200218652-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Kopreski MS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18-JUN-2002
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                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                   Matches
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the atypical endometrial hyperplasia, or Barrett's oesophagus, where the human is at risk for developing a malignancy or premalignancy or is known conclusions at a conclusion of the conclusion
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Peptide products of the human TPC2 and TPC3 gene are involved in regulation of telomere length and activity are useful to diagnose and treat telomere length and activity-related diseases.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TR; telomerase activity-related disease; therapy; fertility; RT-PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human telomerase (hTR) cDNA amplifying R3c downstream RT-PCR primer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 26 BP; 8 A; 6 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Andrews WH, Adams RR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; Pred. No. 59; cive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TCTAACCCTAACTGAGAAGGGCGTAG 70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              rcraacctraacrgagaagggggrag 26
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96US-00583808.
96US-00710249.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAD24246 standard; DNA; 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                07-MAR-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 100.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2002-033174/04.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          telomerase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cancer; pregnancy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (GERO-) GERON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          23-DEC-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13-SEP-1996;
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invention also relates to purified, synthetic or recombinant peptides such as TPC2 or TPC3 used for detecting regulators of telomere length and telomerase activity in mammalian cells and for a variety of related diagnostic and therapeutic purposes. The method is useful for screening, diagnosing, monitoring and treating diseases and other conditions such as ancer, pregnancy, fertility, telomere length and telomerase-activity. The present sequence is a reverse transcription (RT) PCR primer used for amplifying human telomerase (hTR) cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to methods and reagents for regulating telomere length and for modulating telomerase activity in mammalian cells. The invention also relates to purified, synthetic or recombinant peptides such as TPC2 or TPC3 used for detecting regulators of telomere length and telomerase activity in mammalian cells and for a variety of related diagnostic and therapeutic purposes. The method is useful for screening, diagnosting, monitoring and treating diseases and other conditions such as cancer, pregnancy, fertility, telomere length and telomerase-activity. The present sequence is a reverse transcription (RT) PCR primer used for amplifying human telomerase (hTR) CDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Peptide products of the human TPC2 and TPC3 gene are involved in regulation of telomere length and activity are useful to diagnose and treat telomere length and activity-related diseases.
                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; telomerase; TR; telomerase activity-related disease; therapy; cancer; pregnancy; fertility; RT-PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human telomerase (hTR) cDNA amplifying F3b upstream RT-PCR primer.
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100.0%; Pred. No. 59;
iive 0; Mismatches 0; Indels
                                                                                                                                                                                                      Score 26; DB 1; Length 26;
Pred. No. 59;
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                                                                                                                                                                     Sequence 26 BP; 7 A; 3 C; 9 G; 7 T; 0 U; 0 Other;
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                                                                                                                                                                                                                100.0%; Pred. ...
                                                                                                                                                                                                                                                                               145 CTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                                                                        CTTCCACCGTTCATTCTAGAGCAAAC 1
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96US-00583808.
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                                                                                                                                                                                                                                            26; Conservative
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05-JAN-1996;
13-SEP-1996;
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Accurate, reliable diagnosis and/or prognosis of cancer, e.g. breast cancer, by analyzing the RNA components of telomerase in plasma or serum.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   monitoring the evolution of cancers. The method comprises analysing enzyme telomerase RNA in blood plasma or serum. The method is typically used for diagnosing breast, ovarian, stomach or colon cancer and/or monitoring the evolution of the cancers after treatment by chemcherapy or operations. The present sequence is a PCR primer for human telomerape RNA (HTR), which was used in the example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; ss; PCR; telomerase; cancer; pregnancy; fertility; neoplasm; hTR;
RT-PCR; reverse transcriptase PCR; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                          telomerase RNA; PCR primer; cancer; breast; ovarian; stomach; hTR; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to a method for diagnosing and/or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 26 BP; 3 A; 6 C; 11 G; 6 T; 0 U; 0 Other;
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                                                                                                                                                                      Human telomerase RNA, hTR, sense PCR primer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 26;
Pred. No.
70
                       1 rcraacccraacrgagaagggcgrag 26
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45 TCTAACCCTAACTGAGAAGGGCGTAG
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ID ADG82593 standard; DNA; 26
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                                                                                         ABA95496 standard; DNA; 26
                                                                                                                                            (first entry)
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                                                                                                                                                                                                                                                                                                                                                                  (CHEN/) CHEN X Q. (STRO/) STROUN M.
                                                                                                                                                                                                                                                                                                                                                                                               (ANKE/) ANKER P.
                                                                                                                                                                                                                                      Homo sapiens.
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                                                                                                                                            12-MAR-2002
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                                                                                                                 ABA95496;
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                                                                                                                                                                                             Human;
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(ADAM/)

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The invention relates to a monoclonal or isolated polyclonal antibody specifically binds human TPC2 or TPC3 (not defined, appearing as ADG82569 and ADG82571). Also included are an antibody obtained by collecting antiserum from a subject immunised with a peptide comprising at least 10 cantiguous amino acids of TPC-2 pr TPC-3 a host cell secreting the cartibody, a composition for obtaining a TPC2-specific antibody comprising at least 10 contiguous amino acids ADG82569, a composition for obtaining at least 10 contiguous amino acids ADG82569, a composition for obtaining a TPC3-specific antibody comprising at least 10 contiguous amino acids of ADG82571, an isolated, recombinant or synthetic nucleic acid encoding a peptide immunogenic for TPC2-specific antibody is useful for obtaining the nucleic acid. The antibody is useful for determining a condition in a subject associated with a high level of TPC2 or TPC3, and is useful for screening, diagnosing and monitoring conditions auch as cancer, pregnancy or fertility. The antibody is also cueful for treating conditions associated with inappropriate expression of TPC2 or TPC3 cross profile of transcriptase (RT)-PCR primer used to analyse the expression profile of the human telomerase gene, which is co-expressed with TPC-2 and TPC-3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel monoclonal or isolated polyclonal antibody that specifically binds human TPC2 or TPC3. useful for treating neoplasia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Telomerase RNA; TR; promoter; cytotoxin; cancer; neoplasia; hTR; gene therapy; thymidine kinase gene; anticancer therapy; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1; Length 26; 59;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human telomerase RNA gene (hTR) specific primer TRC3R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 26 BP; 8 A; 6 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                         Feng J, Andrews WH, Adams RR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5.8%; Score 26; DB
100.0%; Pred. No. 59;
ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example D; SEQ ID NO 25; 62pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 45 TCTAACCCTAACTGAGAGGGCGTAG 70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 rcraacccraacreagaaggggrag 26
                                                                                                                                   95US-0003492P.
96US-00583808.
96US-00710249.
98US-00220157.
                                                                                          29-JUN-2001; 2001US-00895606
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22-OCT-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 100.
Matches 26; Conservative
                                                                                                                                                                                                                                                      VILLEPONTEAU B.
                                                                                                                                                                                                                                                                              FENG J.
ANDREWS W H.
                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2004-051519/05.
                                                                                                                                                                                                                                                                                                                           (ADAM/) ADAMS R R.
                                                                                                                                                                                                                                                                                                                                                                    Villeponteau B,
US2003207404-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          gene therapy; t
PCR primer; ss.
                                                                                                                                   09-SEP-1995;
05-JAN-1996;
13-SEP-1996;
                                                                                                                                                                                                           23-DEC-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO9938964-A2
                                               06-NOV-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAZ07280;
                                                                                                                                                                                                                                                      (VILL/)
                                                                                                                                                                                                                                                                              (FENG/)
                                                                                                                                                                                                                                                                                                        (ANDR/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            엄
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to a monoclonal or isolated polyclonal antibody specifically binds human TPC2 or TPC3 (not defined, appearing as ADG8256) and ADG82571). Also included are an antibody obtained by collecting a natiserum from a subject immunised with a peptide comprising at least 10 contiguous amino acids of TPC-2 pr TPC-3, a host cell secreting the artibody, a composition for obtaining a TPC3-specific antibody comprising at least 10 contiguous amino acids ADG82569, a composition for obtaining a TPC3-specific antibody comprising a TPC3-specific antibody comprising a TPC3-specific antibody contiguous amino acids of ADG82571, an isolated, recombinant or synthetic nucleic acid encoding a peptide immunogenic for TPC2 specific antibody or TPC3-antibody and a host cell containing the nucleic acid. The antibody is useful for centermining a condition in a subject associated with a high level of TPC2 or TPC3, and is useful for screening, diagnosing and monitoring conditions such as cancer, pregnancy or fertility. The antibody is also useful for treating conditions associated with inappropriate expression of TPC2 or TPC3 such as neoplasia. The present sequence is a reverse transcriptise (RT)-PCR primer used to analyse the expression profile of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; 88; PCR; telomerase; cancer; pregnancy; fertility; neoplasm; hTR; RT-PCR; reverse transcriptase PCR; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel monoclonal or isolated polyclonal antibody that specifically binds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   human telomerase gene, which is co-expressed with TPC-2 and TPC-3.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            human TPC2 or TPC3. useful for treating neoplasia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 26 BP; 7 A; 3 C; 9 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                            Andrews WH, Adams RR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%; Pred. ...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human telomerase gene, hTR, RT-PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       145 CTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example D; SEQ ID NO 26; 62pp; English
                                                                                                                                                                                                      95US-0003492P.
96US-00583808.
96US-00710249.
98US-00220157.
                                                                                                                                                             2001US-00895606
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                         Villeponteau B, Feng J,
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                                                                                                                                                                                                                                                                                                                           VILLEPONTEAU
                                                                                                                                                                                                                                                                                                                                                                      ANDREWS W H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2004-051519/05.
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                                                                                                                                                                                                                                                                                                                                                                                              ADAMS R R.
                                                                  US2003207404-A1.
                       Homo sapiens.
                                                                                                                                                           29-JUN-2001;
                                                                                                                                                                                                                                                                                                                                                 FENG
                                                                                                                                                                                                                                05-JAN-1996;
13-SEP-1996;
                                                                                                                                                                                                                                                                              23-DEC-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       26;
                                                                                                                06-NOV-2003
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                                                                                                                                                                                                                                                                                                                      (VILL/)
(FENG/)
(ANDR/)
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Gaps

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0; Indels

Local Matches

RESULT 77 ADG82592

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RNA (TR) component genes. The TR gene promoter can be linked to a heterologous gene, especially a gene encoding a cytotoxin, for therapy of cancer, especially neoplasias. The telomerase is necessary for the curestricted proliferative capacity of many human cancers. Mutation or unregulation of the telomerase repression pathway may cause reactivation or upregulation of telomerase expression in cancer. Substances, cidentified in the methods, can be used to block transcription from the TR gene promoter through interaction of the 5' regulatory sequences. These cubstances, e.g. antitaense oligonuclectides, transcription factors, peptide nucleic acids and factors that disrupt signal transduction, are useful for cancer therapy. In particular, gene therapy vectors (especially poffs2-codyupp) comprising the promoter and a viral thymidine kinase gene can be used to convert a producy, e.g. gancyclovir, so that cooplasia can be controlled or treated. Direct down-regulation of telomerase RNA gene through manipulation of transcription factors may be effective anticancer therapy and the cloning of the hTR gene promoter allows the analysis of therapeutic molecules which modulate hTR promoter activity. Sequences AAZ01623-80 represents PCR primers for amplifying
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 .;
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                                                                                                                                                                                                                                                                                                                             The invention relates to promoter regions from mouse and human telomerase
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                                                                                                                                                                                                                                     Mouse and human telomerase RNA gene promoters, useful for tumor specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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0
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No. 68;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 25 BP; 6 A; 7 C; 9 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  5.5%; Scor.
100.0%; Pred. No. cc.
... 0; Mismatches
                                                                                                                           (CANC-) CANCER RES CAMPAIGN TECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   175 AATGTCAGCTGCTCGCC 199
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          human TR gene (hTR) promoter sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AATGTCAGCTGCTGGCCCGTTCGCC 1
                                                                                                                                                                                                                                                                                       Disclosure; Fig 6; 109pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Telomerase RNA target sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAA37582 standard; RNA; 25 BP
                                                                                        98GB-00001902.
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                                                    99WO-GB000308
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 100.
Matches 25; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              paternity testing; ss
                                                                                                                                                                                                 WPI; 1999-479183/40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           09-APR-1997;
                                                    29-JAN-1999;
                                                                                        29-JAN-1998;
                                                                                                                                                                                                                                                       gene therapy
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                  05-AUG-1999.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAA37582;
                                                                                                                                                               Keith WN;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      25
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The present sequence represents a human telomerase RNA sequence which is the target of peptide nucleic acid molecules (PNA). The PNA molecules the target of peptide nucleic acid molecules (PNA). The PNA molecules control to mammalian telomerase, and inhibit telomerase activity. Telomerase is a ribonucleoprotein enzyme that telomerase activity. Telomerase is a ribonucleoprotein enzyme that telomerase activity. Telomerase is a ribonucleoprotein enzyme that telomerases contained within the RNA component of the enzyme. The nucleotide sequence contained within the RNA component of the enzyme. The invention relates to PNA molecules having a sequence of no more than 25 bases, which include the sequence GTTAGG. The uncharged nature of the PNA carectases the melting temperature of associating strands, increases the rate of association with targeted nucleic acids, and afforted greater resistance of degradation by proteases or nucleases. The therapeutic PNAs may be used for treating disease conditions such as cancers, neoplasia, hyperplasia, neurodegenerative diseases or nucleases. The immunodeficiancy virus (HIV) infection/AIDS (acquired immunodeficiency syndrome) and associated pathologies, fungal infections, and other cytocoxic or diseases characterized by abnormal telomere metabolism or telomerase activity, in combination with antineoplastic and other cytocoxic or cytostatic agents, antifungal agents, and other nucleotides. PNAs may be used for molecular diagnostics, labelled PNAs are used as hybridization probes to detect or quantitate polymicleotides having a human telomerase and as probes to detect the RNA component of a mammalian telomerase and as inhibitors of telomerase activity. The method of the present invention can be used to detect an immortal or methods of the present invention can be used to detect an immortal or methods of the present invention can be used to detect an immortal or necessaries.
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                                                                                                                                                                       {\it w} peptide nucleic acid (PNA) compounds that inhibit telomerase activity mammalian cells is useful as probes to detect the RNA component of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ;
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                                                                                      Corey DR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1; Length 25;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 25 BP; 7 A; 5 C; 6 G; 0 T; 7 U; 0 Other;
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0
                                                                                   Norton JC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            7; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          68;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 5.5%; Score 25; 72.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       41 TTTGTCTAACCCTAACTGAGAAGGG 65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human telomerase RNA (hTR) strand #1.
                                                                                      Shay JW,
                                                                                                                                                                                                                                                           Example 1; Col 27-28; 45pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAS15453 standard; RNA; 25 BP.
96US-00630019
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                                        (TEXA ) UNIV TEXAS SYSTEM.
                                                                                      Piatyszek MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity 72.0
les 18; Conservative
                                                                                                                                                                            New peptide nucleic acid
                                                                                                                               WPI; 2000-292432/25
09-APR-1996;
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                                                                                      Wright WE,
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Matches
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05-JUN-2000; 2000WO-US015544.
                          99US-0137549P
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                                                                                                                          WPI; 2001-071022/08
                                                     AU J L.
WIENTJES G.
                                                                                               Wientjes G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (GERO-) GERON CORP.
                                                                                                                                                                                    agent to the cell.
                          04-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                08-OCT-1998
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20-DEC-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAV41169;
                                                                                                                                                                                                                                                                                                                                                                                                                                 25
                                                                                                                                                                                                                                                                                                                                              Query Match
                                                     (AUJL/)
(WIEN/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Kim NW,
                                                                                               δu JL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 82
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAV41169/
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                                                                                                                                                                                                                                                                                                                                        The present invention relates to peptide nucleic acids (PNAs), comprising a sequence of 6-25 nucleobases, that inhibit telomerase activity in mammalian cells by hybridising to the RNA component of mammalian telomerase. The PNAs are useful as probes to detect the RNA component of mammalian telomerase and as inhibitors of telomerase activity, or to detect and/or quantitate polymucleotide having the human telomerase RNA component (hTR) sequence, as well as in forensic identification of individuals, such as paternity testing or identification of criminal suspects or unknown descendants based on the HTR gene RRLP pattern. The PNA can be further used for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases. The PNAs in combination with other pharmaccuticals (such as antineoplastic or cytostatic agents) can be used for treating neoplasia, human immunodeficiency virus (HIV) infections, acquired immunodeficiency syndrome (AIDS) and associated pathologies, and other diseases characterised by abnormal telomerase activity. The present sequence representing hTR RNA strand #1 is used to test the inhibition of telomerase activity by the PNAs of the present
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                         New peptide nucleic acids that hybridizes to the RNA component of mammalian telomerase, useful for treating or preventing cancer, inflammation, lymphoprollferative diseases, autoimmune disease, or neurodegenerative diseases.
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                                                                                                                                                                                    Norton JC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 25; DB 1; Length 25;
Pred. No. 68;
7; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Telomerase; cancer; telomere damage; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Seguence 25 BP; 7 A; 5 C; 6 G; 0 T; 7 U; 0 Other;
                                                                                                                                                                                 Piatyszek MA, Corey DR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       41 TTTGTCTAACCCTAACTGAGAAGGG
                                                                                                                                                                                                                                                                                                                  Example 1; Col 29; 46pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human telomerase PCR primer #1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAC93100 standard; DNA; 25 BP
                                                                                   99US-00349532.
                                                                                                           96US-00630019.
97US-00838545.
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                                                                                                                                                     (TEXA ) UNIV TEXAS SYSTEM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            18; Conservative
                                                                                                                                                                                   Shay JW, Wright WE,
                                                                                                                                                                                                              WPI; 2001-638024/73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
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Ното варіелв.
                         US6294650-B1
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                                                                                08-JUL-1999;
                                                                                                           09-APR-1996;
                                                                                                                            09-APR-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14-DEC-2000.
                                                    25-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAC93100;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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                                                                                                                                                                                            The present invention provides a method for inhibiting or reducing the growth of a cell which involves administering to the cell a telomere damage inducing agent and a telomerase inhibitory agent. This can be used in the treatment of aberrant cell growth, including cancers
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Inhibiting or reducing growth of cell for treating cancer, comprising administering telomere damage-inducing agent and telomerase inhibitory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RNA component; human telomerase; antisense oligonucleotide; infection; neuroblastoma; bladder cancer; colon cancer; prostate cancer; cancer; contraception; sterilisation; immunosuppression; therapeutic; hTR; immune system down-regulation; anti-inflammatory therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RNA component of human telomerase (hTR) antisense oligo 14.
                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1; Length 25;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
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                                                                                                                                                                                                                                                                                                                                                                Sequence 25 BP; 4 A; 5 C; 4 G; 12 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 68;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Pred. ...
                                                                                                                                                                                                                                                                                                                                                                                                                             5.5%; Score 25; 00.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            161 TAGAGCAAACAAAAATGTCAGCTG 185
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TAGAGCAAACAAAAATGTCAGCTG 1
                                                                                                                                  Sxample 7; Page 62; 97pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 11; Page 65; 80pp; English
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96US-00770565
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
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Matches 25; Conservative
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inhibitors of telomerase activity, especially in the treatment of cancer
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                                                                                                                                                                                                                       OXCCCCCCCCCCCCCCXRXBAAAXBXBXBXBXBXAAXBXXBXXBXXXXXXXBXAAXBXAAXBX
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antisense oligonucleotides to the RNA component of human telomerase (hTR). These antisense oligonucleotides specifically hybridise to a nucleotide sequence within an accessible region of the hTR, but that does not hybridise to a sequence within the template region of hTR. These oligonucleotides may specifically be used for detection of an RNA component of human telomerase in a sample. This is useful for diagnosing cancer (especially neuroblastona, bladder, colon and prostate cancer).

Can providing prognosis for a cancer patient. The inhibitory accounce. Oligonucleotides can inhibit the telomerase activity level in a cell by interfering with transcription of the RNA component, decreasing the half-cancer of the RNA component into the telomerase activity inhibiting the half-cancer of the RNA component into the telomerase noisonucleotides can be used for inhibiting telomerase activity in both cultured cells and in be used for inhibiting telomerase activity in both cultured cells and in breventing cancer, for contraception or sterilisation, for immunosuppression, and for selectively down-regulating specific branches confine immune system, e.g. a specific subset of T-cells, in anti-cinflammatory therapies or for treating infections by, e.g. yeast,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This is an oligonucleotide used in the measurement of Tm values and their complimentary peptide nucleic acids (PNAs), (e.g. AAT89228-T89233, AAT89243-T89244). PNAs hybridise specifically to an RNA component of mammalian telomerase, and include the sequence GGG for specific hybridisation to the template region of this component. PNAs can be used as probes to detect the RNA component of mammalian telomerase and as
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                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Peptide nucleic acid, PNA; cancer, telomerase, probe, hybridisation, inhibitor, human telomerase RNA, hTR, PCR; oligonucleotide, ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New peptide nucleic acids hybridising to mammalian telomerase RNA - to inhibit telomerase, for treating tumours and other proliferative diseases, also for diagnosis.
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                                                                                                                                                                                                                                                                                                                                     DB 1; Length 30;
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                                                                                                                                                                                                                                                                                                                                                                 1; Indels
                                                                                                                                                                                                                                                                                                          Sequence 30 BP; 4 A; 10 C; 8 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                   5.4%; Score 24.4; D
96.2%; Pred. No. 97;
tive 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                       Piatyszek MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 2; Page 49; 76pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                 Conservative
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                                                                                                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (GERO-) GERON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     09-APR-1996;
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                                                                                                                                                                                                                                                                                                                                                                 25;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                             355
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                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                          56
                                                                                                                                                                                                                                                                                                                                                                 Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               telomerase. Detecting telomerase allows early diagnosis of tumour or infection, before clinical signs manifest. Telomerase inhibitors directed against e.g. Trypanosoma should cause fewer side effects than drugs currently used to treat such infections. The DNA encodes those parts of MTR RNA essential for activity but are significantly shorter than the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cancer; eukaryotic parasite; hTR; vertebrate telomerase; yeast; protozoa; tumour; antibody; polymerase chain reaction; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA encoding essential RNA components of human telomerase - also truncated or recombinant telomerase, useful for diagnosis and treatment of cancer and infection by eukaryotic parasites.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         or to prevent immortalisation of cells in tumours or parasites. The DNA is also used to produce recombinant telomerase, which can then be used conventionally to raise antibodies for diagnostic detection of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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                                                                DB 1; Length 24; 78;
                                                                                                                                            0; Indels
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Sequence 24 BP; 7 A; 5 C; 5 G; 7 T; 0 U; 0 Other;
                                                     Query Match
Best Local Similarity 100.0%; Pred. No. 'o,
Matches 24; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human telomerase template PCR primer TRC31.
                                                                                                                                                                                                                    41 TITGICIAACCCIAACTGAGAAGG 64
                                                                                                                                                                                                                                                                                    1 rrrgrcraacccraacrgagagg 24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (COLD-) COLD SPRING HARBOR LAB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAT58807 standard; DNA; 24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAT58807;
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primers AAV68464-65 are used for the RT-PCR amplification of the RNA component of human telomerase (hTR). The invention relates to an activate complex (A1) that comprises: (a) an antisense oligonuclectide, complementary to a 12-25 nucleotide portion of hTR, with a hydroxyl moiety at the first end; and (b) a linker attached to the first end, and (c) an activator of RNase L attached to the linker. The citrator-antisense complex may be used for inhibiting the growth of a telomerase-expressing malignant cell or tumour. The complex is used to specifically cleave the ribonuclectide portion of a telomerase enzyme. The complex inhibiting growth of telomerase expressing malignant cells from brain tumour malignant glioma, breast tumour, renal cell cancer. The complex inhibiting spowth of telomerase expressing malignant cells from brain tumour malignant glioma, breast tumour, renal cell cancer. Celisorders, may be treated including various acute and chronic leukemias, polycthemia vera, Hodgkin's and non-Hodgkin's lymphomas, multiple myeaven.
                                                                                                                                                                                                                                                                                                                     cleave; brain; tumour malignant glioma; breast tumour; renal cell cancer; melanoma; prostete cancer; leukemia; polytchemia vera; myelloma; sarcoma; Hodgkin's lymphoma; Waldenstrom's macroglobulinemia; heavy chain disease; carcinoma; chemotherapeutic; antisense; RT-PCR; primer; ss.
                                                                                                                                                                                                                                                                                                    Human; telomerase; hTR; activator-antisense complex; malignant; enzyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New RNase L activator-telomerase antisense complex - useful to inhibit telomerase activity in telomerase-expressing malignancies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               tumours, including numerous sarcomas and carcinomas. The complex is preferably administered in combination with a chemotherapeutic agent, particularly either cisplatin, doxorubicin, mitomycin, daunorubicin,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Torrence
                                                                                                                                                                                                                                                            Human telomerase RNA (hTR) amplifying RT-PCR primer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              bleomycin, actinomycin D, or neocarzinostatin
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CTCTGTCAGCCGCGGTCTCTCGG 334
                                   24 crcrercaecececesererese 1
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                                                                                                                                       AAV68465 standard; DNA; 24 BP.
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                                                                                                                                                                                                                  (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1998-609972/51.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (CLEV-) CLEVELAND
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                                                                                                                                                                                                                                                                                                                                                                                                                                                Ношо варіепв
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             03-FEB-1998;
                                                                                                                                                                                                                     22-MAR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         29-OCT-1998
                                                                                                                                                                               AAV68465;
                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic
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Component of human telomerase (hTR). The invention relates to an activator-antisense complex that comprises: (a) an antisense to an activator-antisense complex that comprises: (a) an antisense to an object of complementary to a 12-25 nucleotide portion of hTR, with a hydroxyl moiety at the first end; and (b) a linker attached to the linker. The cityst end, and activator of RNase L attached to the linker. The city activator-antisense complex may be used for inhibiting the growth of a telomerase expressing malignant cell or tumour. The complex is used to specifically cleave the ribonucleotide portion of a telomerase enzyme. The complex inhibits growth of telomerase expressing malignant cells from brain tumour malignant glloma, breast tumour, renal cell cancer, melanoma, and prostate cancer. Many other malignancies and related disorders, may be treated including various acute and chronic leukemias, polycthemia vera, Hodgkin's and non-Hodgkin's lymphomas, multiple tumours, including numerous sarcomas and carcinomas. The complex is tumours, including numerous sarcomas and carcinomas. The complex is treated in combination with a chemotherapeutic agent,
                                                                                                                                                                                                                                                        Human; telomerase; hTR; activator-antisense complex; malignant; enzyme; cleave; brain; tumour malignant glioma; breast tumour; renal cancer; melanoma; prostate cancer; leukemia; polycthemia vera; myeloma; sarcoma; Hodgkin's lymphoma; Waldenstrom's macroglobulinemia; heavy chain disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New RNase L activator-telomerase antisense complex - useful to inhibit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       particularly either cisplatin, doxorubicin, mitomycin, daunorubicin,
bleomycin, actinomycin D, or neocarzinostatin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                                                                                                                                                                  carcinoma; chemotherapeutic; antisense; RT-PCR; primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         telomerase activity in telomerase-expressing malignancies.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Torrence PF;
                                                                                                                                                                                                                      Human telomerase RNA (hTR) amplifying RT-PCR primer.
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CGTGCACCCAGGACTCGGCTCACA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            41 TTTGTCTAACCCTAACTGAGAAGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example; Page 41; 81pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (CLEV-) CLEVELAND CLINIC FOUND. (USSH ) US NAT INST OF HEALTH.
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98US-00018125.
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                                                                                                              AAV68464 standard; DNA; 24
                                                                                                                                                                                    22-MAR-1999 (first entry)
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1es 24; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1998-609972/51.
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                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
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03-FEB-1998;
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Query Match Best Local S

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Score 24; DB 1; Length 24; Pred. No. 78;

AAZ07273/c

RESULT 87

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AAZ07279 standard; DNA; 24 BP
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                                                  22-OCT-1999 (first entry)
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Best Local Similarity
                                                                                                                                                                  PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                     39-JAN-1998;
                                                                                                                                                                                                                    Homo sapiens
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                 AAZ07263;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to promoter regions from mouse and numerated to a hereacologous gene, especially a gene promoter can be linked to a cherologous gene, especially a gene encoding a cytotoxin, for therapy of cancer, especially neoplasias. The telomerase is necessary for the cancer, especially neoplasias. The telomerase is necessary for the unrestricted proliferative capacity of many human cancers. Mutation or dysregulation of the telomerase expression in cancer. Substances, identified in the methods, can be used to block transcription from the TR gene promoter through interaction of the 5' regulatory sequences. These substances, e.g. antisense oligonucleotides, transcription factors, or useful for cancer therapy. In particular, gene therapy vectors useful for cancer therapy. In particular, gene therapy vectors (especially pGT62-codAupp) comprising the promoter and a viral thymidine complasia can be convert a prodrug, e.g. gancyclovir, so that neoplasia can be convert a prodrug, e.g. gancyclovir, so that comparase RNA gene through manipulation of transcription factors may be effective anticancer therapy and the cloning of the HTR gene promoter allows the analysis of therapeutic molecules which modulate hTR promoter activity. Sequences AAZ07623-80 represents PCR primers for amplifying thuman TR gene (HTR) promoter sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to promoter regions from mouse and human telomerase
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                                                                                                                                                                                                                      Telomerase RNA; TR; promoter; cytotoxin; cancer; neoplasia; hTR; gene therapy; thymidine kinase gene; anticancer therapy; human;
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100.0%; Pred. No. 78;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                     Human telomerase RNA gene (hTR) specific primer hTR5.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (CANC-) CANCER RES CAMPAIGN TECHNOLOGY.
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                                                                         AAZ07273 standard; DNA; 24 BP.
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24; Conservative
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                                                                                                                                                                                                                                                                 PCR primer; 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         gene therapy.
                                                                                                                                                                                                                                                                                                                       Homo sapiens
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AAZ07263/c ID AAZ07263 standard; DNA; 24 BP.

RESULT 88

Matches

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NAM (TR) component genes. The TR gene promoter can be linked to a checrologous gene, especially a gene promoter can be linked to a checrologous gene, especially a gene promoter can be linked to a cancer. Superially neoplasias. The telomerase is necessary for the cancer is especially neoplasias. The telomerase is necessary for the currestricted proliferative capacity of many human cancers. Mutation or dysregulation of the telomerase repression pathway may cause reactivation or the methods, can be used to block transcription from the TR gene promoter through interaction of the 5' regulatory sequences. These substances, e.g. antisense oligomucleotides, transcription factors, peptide nucleic acids and factors that disrupt signal transduction, are useful for cancer therapy. In particular, gene therapy vectors (especially pGT62-codAupp) comprising the promoter and a viral thymidine consecting and be used to convert a producy. Generally pGT62-codAupp) comprising the promoter and a viral thymidine concortive anticancer therapy and the cloning of the franscription factors may be effective anticancer therapy and the coloning of the franscription actions activity. Sequences AAZ07623-80 represents PCR primers for amplifying thuman TR gene (hTR) promoter sequence
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                                                                                Telomerase RNA; TR; promoter; cytotoxin; cancer; neoplasia; hTR; gene therapy; thymidine kinase gene; anticancer therapy; human;
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Human telomerase RNA gene (hTR) specific primer hTR5.
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RNA (TR) component genes. The TR gene promoter can be linked to a

Cancer, especially agene encoding a cytotoxin, for therapy of

cancer, especially neoplasias. The telomerase is necessary for the

cancer, especially neoplasias. The telomerase is necessary for the

curregulation of the telomerase repression pathway may cause reactivation of

cypergulation of the telomerase expression in cancer. Substances,

clantified in the methods, can be used to block transcription from the TR

cyperide promoter through interaction of the 5' regulatory sequences. These

substances, e.g. antisense oligonucleotides, transcription factors,

cyperide nucleic acids and factors that disrupt signal transduction, are

useful for cancer therapy. In particular, gene therapy vectors

chepcially pG762-codAupp) comprising the promoter and a viral thymidine

ckinase gene can be used to convert a prodrug, e.g. gancyclovir, so that

celomerase RNA gene through manipulation of transcription factors may be

effective anticancer therapy and the cloning of the hTR gene promoter

allows the analysis of therapeutic molecules which modulate hTR promoter

callows the analysis of therapeutic molecules which modulate hTR promoter

burner man and a prodrugh of the pTR promoter allows the analysis of the pTR persents pCR primers for amplifying
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                                                                                                                                                                                                                                                                                                          Mouse and human telomerase RNA gene promoters, useful for tumor specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                        nerase RNA; TR; promoter; cytotoxin; cancer; neoplasia; hTR; therapy; thymidine kinase gene; anticancer therapy; human;
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Human telomerase RNA gene (hTR) specific primer TRC3F.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 24 BP; 8 A; 6 C; 6 G; 4 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TR gene (hTR) promoter sequence
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                           relomerase RNA; TR;
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                                                                                           Homo sapiens.
                                                                                                                    WO9938964-A2
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                                                                               Synthetic
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The present sequence represents a peptide nucleic acid molecule which hybridises to the mRNA component of mammalian telomerase, and inhibits telomerase activity. Telomerase is a ribbonucleoprotein enzyme that synthesizes one strand of the telomeric DNA, using as a template an 11 nucleotide sequence contained within the RNA component of the enzyme. The invention relates to PNA molecules having a sequence of no more than 25 concess, which include the sequence GTTAGG. The uncharged nature of the PNA backbone increases the melting temperature of associating strands, increases the rate of association with targeted nucleic acids, and affords greater resistance of degradation by proteases or nucleases. The therapeutic PNAs may be used for treating disease conditions such as cancers, neoplasia, hyperplasia, neurodegenerature diseases. The ferminal strands and associated pathologies, fungal infections, and other conditions and other cytostatic agents, antifungal agents, fungal infections, and other diseases characterized by abnormal telomere metabolism or telomerase activity, in combination with antineoplastic and other cytotoxic or cytostatic agents, antifungal agents, and other nucleotides. PNAs may be used for molecular diagnostics, labelled PNAs are used as Hybridization probes to detect or quantitate polynucleotides having a human telomerase and as individuals, e.g. paternity testing, based on hTR gene restriction of individuals, e.g. paternity testing, based on hTR gene restriction fragment length polymorphism (RFLP) pattern. PNAs are also useful as probes to detect the RNA component of a mammalian telomerase and as inhibitors of telomerase activity. The method of the present invention and person inhibitors of telomerase activity. The method of detected as cancerous conditions to be detected with increased confidence and methods of the present invention can be used to detect an immortal or methods of the present invention acells are detected an immortal or cell expresses telomerase activity and its RNA component
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New peptide nucleic acid (PNA) compounds that inhibit telomerase activity in mammalian cells is useful as probes to detect the RNA component of a mammalian telomerase.
                                                                                                                              /*tag= a /*tag= a /*tag= a /*tag= nucleic acid molecule, where N-(2-aminoethyl)glycine units are linked to nucleotide bases via glycine amino N through a methylenecarbonyl linker.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gape
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Corey DR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1; Length 24; 78;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 24 BP; 7 A; 5 C; 5 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          .
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           5.3%; Score 24;
100.0%; Pred. No.
:ive 0; Mismatc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TTTGTCTAACCCTAACTGAGAAGG 24
                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 2; Col 33; 45pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          97US-00838545.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 96US-00630019
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Piatyszek MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (TEXA ) UNIV TEXAS SYSTEM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2000-292432/25.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
les 24; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          09-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 09-APR-1996;
                                                                           Key
misc feature
                                                                                                                                                                                                                                                                                                                                   JS6046307-A
                                                                                                                                                                                                                                                                                                                                                                                                               04-APR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Wright WE,
Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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ADQ36831 standard; DNA; 24 BP.
                                                       ADQ36831;
RESULT 92
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               ADQ3683
                                          8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention relates to peptide nucleic acids (DNAs), comprising a sequence of 6-25 nucleobases, that inhibit telomerase activity in mammalian cells by hybridising to the RNA component of mammalian cells by hybridising to the RNA component of mammalian telomerase and as inhibitors of telomerase activity, or to detect and/or quantitate polynucleotide having the human telomerase RNA component (hTR) sequence, as well as in forensic identification of individuals, such as paternity testing or identification of criminal suspects or unknown descendants based on the HTR gene RFLP pattern. The PNA can be further used for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases. The PNAs in combination with other pharmaceuticals (such as antineoplastic or cytostatic agents) can be used for treating neoplasia, human immunodeficiency virus (HIV) infections, acquired immunodeficiency syndrome (AIDS) and associated pathologies, and other diseases characterised by abnormal telomera metabolism or telomerase activity. The present sequence representing a DNA oligonucleotide is complementary to some of the PNAs of the present invention, and is used
                                                                                                                                                          hTR gene RFLP pattern; cancer; inflammation; forensic; lymphoproliferative disease; autoimmune disease; hyperplasia; neurodegenerative disease; neoplasia; HIV; AIDS; cytostatic; human immunodeficiency virus; acquired immunodeficiency syndrome; telomere metabolism; anti-inflammatory; immunosuppressive; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New peptide nucleic acids that hybridizes to the RNA component of mammalian telomerase, useful for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, on neurodegenerative diseases.
                                                                                                               Oligonucleotide #1 used in melting temperature studies of PNAs.
                                                                                                                                          paternity testing; human telomerase RNA component;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Norton JC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 24; DB 1; Length 24; Pred. No. 78;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Seguence 24 BP; 7 A; 5 C; 5 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Corey DR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; Pred. w..
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Piatyszek MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 2; Col 34; 46pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               melting temperature studies
                           AAS15445 standard; DNA; 24 BP.
                                                                                                                                                                                                                                                                                                                                                                             96US-00630019.
97US-00838545.
                                                                                                                                                                                                                                                                                                                                                99US-00349532
                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                       (TEXA ) UNIV TEXAS SYSTEM
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Shay JW, Wright WE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2001-638024/73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                        US6294650-B1
                                                                                                                                                                                                                                                                                                                                                08-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                             09-APR-1996;
09-APR-1997;
                                                                                   14-FEB-2002
                                                                                                                                                                                                                                                                                                                     25-SEP-2001
                                                                                                                                            Mammalian;
                                                                                                                                                                                                                                                               Synthetic.
                                                       AAS15445;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
RESULT 91
            AAS15445
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Provided are an Adenovirus vector with a telomerase promoter associated with cancer occurrence and a method for using the same for anti-cancer treatment. An Adenovirus expression vector ps-HTERT-TK is produced by using an expression cassette consisting of suicide gene HSV-TK, as late plyadenylation signal for human telomerase hTERT promoter and bovine growth hormone (BGH). Adenovirus clone Ad-hT-TK(KCCM 10387) is used for anti-cancer gene therapy. The present sequence represents a primer of the
                                                                                Adenovirus vector; cancer; anti-cancer; pS-hTERT-TK; suicide gene; HSV-TK; human telomerase; hTERT promoter; bovine growth hormone; BGH; Adenovirus clone; Ad-hT-TK; Cytostatic; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New adenovirus vector comprising telomerase promoter, useful in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Telomerase, body fluid; cancer, tumour; screening; TRAP; diagnosis; telomeric repeat amplification protocol; detection; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         5.3%; Score 24; DB 1; Length 24;
100.0%; Pred. No. 78;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 24 BP; 7 A; 5 C; 5 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human telomerase RNA template PCR primer R3C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%; Prec. ...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; SEQ ID NO 3; 1pp; Korean.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 41 TTTGTCTAACCCTAACTGAGAAGG 64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 TTTGTCTAACCCTAACTGAGAAGG 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP.
                                                                                                                                                                                                                                                                                                          27-JUN-2002; 2002KR-00038076.
                                                                                                                                                                                                                                                                                                                                                    27-JUN-2002; 2002KR-00038076.
                                           Primer of the invention #3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAZ08704 standard; DNA; 25
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20-OCT-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2004-363161/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            treatment of cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Kim YT, Song JS;
                                                                                                                                                                                                                                                                                                                                                                                                                    SONG J S.
                                                                                                                                                                                                                                                                                                                                                                                                  KIM Y T.
                                                                                                                                                                                                                     KR2004002322-A.
                                                                                                                                                                          Unidentified,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO9941406-A1.
26-AUG-2004
                                                                                                                                                                                                                                                               07-JAN-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           nvention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                  (KIMY/)
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Gaps

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0; Indels

Conservative

24;

Matches

41

g ð

64

TTTGTCTAACCCTAACTGAGAAGG 24

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New peptide
to inhibit t
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                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                              RESULT 95
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                                                                                                                                                                                                                                                                                                                                                            AAA37568
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                                                                                                                                                                                                                                 cellular portion of body fluid from a cancer patient using a modified cellular portion of body fluid from a cancer patient using a modified tellular portion of body fluid from a cancer patient using a modified cellular protecol (TRAP). A method for detecting cancer comprises: (a) removing the cellular portion of a body fluid specimen from the patient; (b) preparing a protein extract from the body fluid remainder; (c) assaying the extract for the presence and quantity of telomerase RNA or telomerase activity; and (d) comparing the results with normal levels, to determine the presence of cancer. The methods are used in cancer diagnosis and prognosis, and also to monitor cancer therapy effectiveness. Unlike prior art telomerase activity assays in cancer patients, the method allows noninvasive sample collection. The methods are also more reliable and less tumour specific than other methods which detect circulating tumour markers. The present sequence represents a human telomerase RNA template PCR primer used in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                    Detecting telomerase activity in non-cellular body fluid using a modified telomeric repeat amplification protocol.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Peptide nucleic acid; PNA; cancer; telomerase; probe; hybridisation; inhibitor; human telomerase RNA; hTR; PCR; oligonucleotide; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA oligonucleotide 2, used in the measurement of Tm values.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 24; DB 1; Length 25;
Pred. No. 82;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Norton JC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Indels
                                                                                                                  Abruzzo LV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 25 BP; 7 A; 3 C; 9 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Corey D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      5.3%; Scor.
100.0%; Pred. No. c...
0; Mismatches
                                                                                                                  Highsmith E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CTTCCACCGTTCATTCTAGAGCAA 168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Piatyszek MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            N
                                                                                                                                                                                                             Disclosure, Page 16; 32pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CTTCCACCGTTCATTCTAGAGCAA
                                                                                        (UYMA-) UNIV MARYLAND BALTIMORE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          97WO-US005931
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                                      99WO-US003302
                                                               98US-0074793P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAT89246 standard; DNA; 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                  Stamberg J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Shay JW, Wright WE,
                                                                                                                                              WPI; 1999-508655/42.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1997-512647/47.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                      16-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO9738013-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          09-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    09-APR-1996;
                                                                                                                  Strovel JW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12-MAY-1998
             19-AUG-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16-OCT-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    145
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAT89246;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matchee
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Peptide nucleic acid; PNA; telomerase; ribonucleoprotein enzyme; cancer; inhibitor; neoplasia; neurodegenerative disease; aging; hyperplasia; AIDS; HIV; fungal infection; forensic identification; detect; tumour; paternity testing; ss.
need
                                                                                            This is an oligonucleotide used in the measurement of Tm values and thei complimentary peptide nucleic acids (PNAs), (e.g. AAT89225-T89227). PNAs hybridise specifically to an RNA component of mammalian telomerase, and include the sequence GGG for specific hybridisation to the template region of this component. PNAs can be used as probes to detect the RNA component of mammalian telomerase and as inhibitors of telomerase activity, especially in the treatment of cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note= "Peptide nucleic acid molecule, where N-(2-aminoethyl)glycine units are linked to nucleotide bases via glycine amino N through a methylenecarbonyl linker"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present sequence represents a peptide nucleic acid molecule which hybridises to the mRNA component of mammalian telomerase, and inhibits telomerase activity. Telomerase is a ribonucleoprotein enzyme that
                                                                                                                                                                                                                                                                                                  Gaps
nucleic acids hybridising to mammalian telomerase RNA -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DR;
                                                                                                                                                                                                                                                               DB 1; Length 23;
89;
                                                                                                                                                                                                                                                                                                Indele
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PNA sequence #26 used to inhibit telomerase activity.
                                                                                                                                                                                                                                  Sequence 23 BP; 5 A; 7 C; 1 G; 10 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Norton JC,
                                                                                                                                                                                                                                                             5.1%; Score 23; DB
100.0%; Pred. No. 89;
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Shay JW,
                                                                                                                                                                                                                                                                                                                                   57
                                                                                                                                                                                                                                                                                                                                                                  23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
                                                                Example 2; Page 49; 76pp; English
                                                                                                                                                                                                                                                                                                                                 CCATTTTTGTCTAACCCTAACT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 2; Col 33; 45pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      ВP
                                 diseases, also for diagnosis.
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             telomerase, for
                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAA37568 standard; DNA; 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Piatyszek MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (TEXA ) UNIV TEXAS SYSTEM
                                                                                                                                                                                                                                                                                                23; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               mammalian telomerase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2000-292432/25
                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           39-APR-1996;
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controlled sequence contained within the RNA component of the enzyme. The nucleotide sequence contained within the RNA component of the enzyme. The nucleotide sequence of no more than 25 bases, which include the sequence of TayAGG. The uncharged nature of the PNA controlled the sequence of associating strands, and affords greater realstance of degradation with targeted nucleia acids, and affords greater realstance of degradation by proteases or nucleases. The therapeutic PNAs may be used for treating disease conditions such as the cancers, neoplasia, hyperplasia, neurodegenerative diseases. The immunodeficiency virus (HVV) infection/AIDS (acquired immunodeficiency syndrome) and associated pathologies, fungal infections, and other cyndrome) and associated by abnormal telomere metabolism or telomerase activity, in combination with antineoplastic and other cytotoxic or cytostatic agents, antifungal agents, and other nucleotides. PNAs may be used for molecular disagnostics, labelled PNAs are used as hybridization cytostatic agents, antifungal agents, and other nucleotides. PNAs may be used for molecular disagnostics, labelled PNAs are used as hybridization or probes to detect or quantitate polynucleotides having a human telomerase of individuals, e.g. paternity testing, based on HTR sequence. PNA propose so used for forensic identification of inhibitors of telomerase activity. The membrdo of the present inventions to be detected with increased confidence and so probbs to detect the RNA component of a mammalian telomerase and as inhibitors of telomerase activity. The membrdo of the present invention can be used to detect an immortal or methods of the present invention can be used to detect an immortal or enclopastic cell expresses telomerase activity and its RNA component
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      hTR gene RFLP pattern; cancer; inflammation; forensic; lymphoproliferative disease; autoimmune disease; hyperplasia; lymphoproliferative disease; neoplasia; HIV; AIDS; cytostatic; human immunodeficiency virus; acquired immunodeficiency syndrome; telomere metabolism; anti-inflammatory; immunosuppressive; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Oligonucleotide #2 used in melting temperature studies of PNAs.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 23; DB 1; Length 23;
Pred. No. 89;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  5.1%; Scor.
100.0%; Pre
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97US-00838545.
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Matches
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Norton JC;

Shay JW, Wright WE, Piatyszek MA, Corey DR,

WPI; 2001-638024/73

(TEXA) UNIV TEXAS SYSTEM

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The present invention relates to peptide nucleic acids (PNAs), comprising a sequence of 6-25 nucleobases, that inhibit telomerase activity in mammalian cells by hybridishing to the RNA component of mammalian cells by hybridishing to the RNA component of mammalian telomerase and as inhibitors of telomerase activity, or to detect and/or quantitate polynucleotide having the human telomerase RNA component (hTR) sequence, as well as in forensic identification of individuals, such as paternity testing or identification of criminal suspects or unknown descendants based on the hTR gene RRLP pattern. The Suspects or unknown descendants based on the hTR gene RRLP pattern. The suspects or unknown descendants based on the hTR gene RRLP pattern. The Suspects or unknown descendants have not identification of criminal suspects or unknown descendants have not be need for treating continuous lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases, the PRAs in combination with other pharmaceuticals (such as antimunodeficiency virus (HIV) infections, acquired immunodeficiency virus (HIV) infections, acquired immunodeficiency syndrome (AIDS) and associated pathologies, and other diseases characterised by abnormal telomere metabolism or telomerase activity. The present sequence repersenting a DNA oligonucleotide is complementary to some of the PNAs of the present invention, and is used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; double-stranded RNA; micro-RNA; mirNA; short interfering siRNA; shrNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
         New peptide nucleic acids that hybridizes to the RNA component of mammalian telomerase, useful for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases.
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89;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human TERC mRNA transcript target sequence, SEQ ID 521.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 23 BP; 5 A; 7 C; 1 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity 100.0%; Pred. No. 89; es 23; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              57
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 ccarrrrrgrcraacccraacr 23
                                                                                                                       Example 2; Col 34; 46pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              35 CCATTTTTGTCTAACCCTAACT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       n melting temperature studies
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2002US-0406784P.
2002US-0408378P.
2002US-0409293P.
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2002US-0386782P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           5.1%;
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05-SEP-2002;
09-SEP-2002;
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Matches
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New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.

11-MAR-2002; 2002US-0363124P. 06-UTV-2002; 2002US-03867B2P. 17-UTV-2002; 2002US-0396600P. 29-AUG-2002; 2002US-0406784P. 05-SEP-2002; 2002US-040837BP. 05-SEP-2002; 2002US-040939P. 15-JAN-2003; 2003US-0440129P.

(RIBO-) RIBOZYME PHARM INC. Mcswiggen J, Beigelman L;

WPI; 2003-689777/65.

11-FEB-2003; 2003WO-US004088

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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and antiense regions, or alternatively are assembled from a sense oligonuclectide and an antisense oligonuclectide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonuclectides, and can be chemically synthesised, can contain deoxyribonuclectides, and can be chemically synthesised, captessed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA, conjugates and/or complexes of siNA, and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single engine plants). The present sequence contains and persons and persons and proper contains and persons ag
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression medulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
                                                                                                                                                           New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      mapping (e.g., of single nucleotide polymorphisms). represents a human TERC transcript target sequence.
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                                                                                                                                                                                                                           Example 3; SEQ ID NO 521; 145pp; English.
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15-JAN-2003; 2003US-0440129P.
                                        (RIBO-) RIBOZYME PHARM INC
                                                                                 Mcswiggen J, Beigelman L;
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                                                                                                                         WPI; 2003-689777/65.
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise sense and antisense of adouble or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double stranded RNA, micro-RNA (miRNA) and short can antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA). The siNAs and be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treatment of a variety of conditions. They may be used for treatment rejection, or autoimmune or age-related diseases, ce.g. multiple sclerosis, lupus erythematosus, AlDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening diagnosis, lupus erythematosus, AlDS, macular degeneration, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence mapping (e.g., of single nucleotide polymorphisms). The present sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human TERC mRNA transcript target sequence, SEQ ID 522.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 23 BP; 6 A; 10 C; 6 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pred. No. 89;
Disclosure; SEQ ID NO 530; 145pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           5.1%; Score 23;
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Best Local Similarity 95.77
Matches 22, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADF93795;
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense coupling stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and antisense oligonucleotide. Specifically, the siNAs include short that repring RNA (siNRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (shRNA). The siNAs can be unmodified or chemically spothesised, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, can complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treatsing cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AlDS, macular degeneration, stranglanting diagnosis, therapeutic target identification and validation, sortening diagnosis, therapeutic target identification and validation, sortening diagnosis, therapeutic target identification and validation, sortening diagnosis, therapeutic target identification and validation, and mense of entities and rheumatoid arthritis. The siNAs are also useful for drug screening cancer, and service described and remaining diagnosis, therapeutic target identification and validation, and mense of entities and rheumatoid arthritis. The siNAs are also useful for drug screening cancer, and the sinas are also useful described diseases.
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drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 3; SEQ ID NO 522; 145pp; English.
                                                                                                                                                                                                                                                                                                                                2002US-0386782P.
2002US-0396600P.
2002US-0406784P.
2002US-0408378P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Beigelman L;
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                                                                                                                                 WO2003070742-A1.
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06-JUN-2002;
17-JUL-2002;
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05-SEP-2002;
09-SEP-2002;
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                                                                                Homo sapiens
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o; Gaps . 0 5.1%; Score 23; DB 1; Length 23; 78.3%; Pred. No. 89; 0; Indels ve 5; Mismatches 0; Indels 298 GCGAAGAGTTGGGCTCTGTCAGC 320 78.3%; ADF93793 standard; mRNA; 23 Conservative Best Local Similarity 18; Query Match RESULT 100 Matches ADF93793 DP 12225 ઠે

(first entry)

26-FEB-2004

ADF93793;

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Cytostatic, vasotropic, protozoacide, immunosuppressive, dermatological; neuroprotective, anti-HIV; ophthalmological; antiulcer; anti-rheumatic;
                                                antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; micro-RNA; micro-RNA; micro-RNA; micro-RNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis, therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
                                                                                                                                                                                                                                                                                                                                                                                                 New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
        Human TERC mRNA transcript target sequence, SEQ ID 520.
                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 3; SEQ ID NO 520; 145pp; English.
                                                                                                                                                                                                                                                                                   05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409293P.
15-JAN-2003; 2003US-0440129P.
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                                                                                                                                                                                                                                                                        2002US-0406784P.
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                                                                                                                                                         WO2003070742-A1
                                                                                                                                                                                                                                                                                                                                                         Mcswiggen J,
                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                           17-JUL-2002;
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The invention relates to short interfering nucleic acids (siNA) which compures the synession of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense or alternatively are assembled from a sense oligonucleotide and antisense cligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double stranded RNA, micro-RNA (miRNA) and short can contain deoxyribonucleotides, and can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised.

CC expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue cyprossion of the telomerase genes in cells, tissue cyprossion of the telomerase spens in cells, tissue cyprospants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, lupus erythematosus, ALDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, lupus erythematomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence cypenesents a human TERC transcript target sequence. Sequence 23 BP; 4 A; 9 C; 3 G; 0 T; 7 U; 0 Other;

Gaps .. DB 1; Length 23; 0; Indels Score 23; DB 1; Pred. No. 89; 7; Mismatches 144 CCTTCCACGTTCATTCTAGAGC 166 23 1 ccuuccaccguucauucuagagc Query Match 5.1%; Best Local Similarity 69.6%; Matches 16; Conservative 8 g

Sequence 23 BP; 2 A; 11 C; 4 G; 0 T; 6 U; 0 Other;

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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonuclectide and an tisense oligonuclectide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (shRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonuclectides, and can be chemically synthesised. Can contain deoxyribonuclectides, and can be chemically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express ginA. The siNAs are used to modilate expression of the telomerase genes in cells, tissue cased to modilate expression of the telomerase genes in cells, tissue caplants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically processed), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosuus, AlbS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug secenting, diagnosis, therapeutic target identification and validation, apping (e.g., of single nucleotide polymorphisms). The present sequence mapping (e.g., of single nucleotide polymorphisms). The present sequence
                                                                                                                                                                                                              Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; slNA; short interfering nucleic acid; slNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
                                                                                                                                                                          Human TERT mRNA transcript target sequence, SEQ ID 529.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; SEQ ID NO 529; 145pp; English.
                                           ADF93802 standard; mRNA; 23 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2002US-0396600P.
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                                                                                                                              (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
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                                                                                                                              26-FEB-2004
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                                                                                     ADF93802;
RESULT 101
                      ADF93802
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonuclectide and an intense oligonuclectide. Specifically, the siNAs include short interfering RNA (siRNA), double-strands RNA, micro-RNA (miRNA) and short interfering RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonuclectides, and can be chemically synthesised. Cexpressed from a vector or erzymatically synthesised. The invention also relates to kits for the in vitro of all very of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be
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                                           Gaps
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     Score 23; DB 1; Length 23; Pred. No. 89;
                                           Indels
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                                           6; Mismatches
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   5.1%; Score 23; 73.9%; Pred. No.
                                                                              136 GCCTGCCGCTTCCACCGTTCAT 158
                                                                                                    1 GCCUGCCGCCUUCCACCGUUCAU 23
                                                                                                                                                                                                              ADF93804 standard; mRNA; 23 BP.
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2002US-0386782P.
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Query Match
Best Local Similarity 73.9°
Matches 17; Conservative
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                                                                                                                                                                        RESULT 102
ADF93804
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                 protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents a human TERT transcript target sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cytostatic, vasotropic, protozoacide, immunosuppressive, dermatological, neuroprotective, anti-HIV; ophthalmological, antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shray, expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
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   treating cancer, restenosis, infectious diseases (specifically
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                                                                                                                                                                Sequence 23 BP; 2 A; 8 C; 9 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                           4; Mismatches
                                                                                                                                                                                                                           89;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; SEQ ID NO 528; 145pp; English.
                                                                                                                                                                                                        Score 23;
Pred. No.
                                                                                                                                                                                                                                                                              395 GCGCGCGCGATTCCCTGAGCTG 417
                                                                                                                                                                                                                                                                                                   1 GCGCGCGCGAUCCCCUGAGCUG 23
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                                                                                                                                                                                                    5.1%;
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2002US-0386782P.
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2002US-0406784P.
2002US-0408378P.
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                                                                                                                                                                                                Query Match
Best Local Similarity 82.6<sup>5</sup>
Matches 19, Conservative
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29-AUG-2002;
05-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-FEB-2002;
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06-JUN-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  26-FEB-2004
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 used for
                                                                                                                                                                                                                                                                                                                                                                                                                                                ADF93801;
                                                                                                                                                                                                                                                                                                                                                                          RESULT 103
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interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (shRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA, conjugates and/or complexes of siNA, and vectors that express siNA. The siNas are used to modulate expression of the telomerase genes in cells, tissue captains or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restencisis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single mucleotide polymorphisms). The present sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New double-stranded short interfering nucleic acid molecule, useful for down-regulating the expression of an endogenous mammalian target gene or for treating diseases that respond to modulation of gene expression or activity.
                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       double-stranded short interfering nucleic acid; siNA; antibarkinsonian; antiarteriosclerotic; neuroprotective; nootropic; antiparkinsonian; anticonvulsant; pulmonary disease; restenosis; atherosclerosis; Alzheimer's; Parkinson's; epilepsy; dementia; huntington's; amyotrophic lateral sclerosis; gene therapy; target; ss; hTR.
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                                                                                                                                                                                                                                                                                                                                                           DB 1; Length 23;
                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                      Sequence 23 BP; 2 A; 3 C; 14 G; 0 T; 4 U; 0 Other;
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0
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                                                                                                                                                                                                                                                                                                                                                                                            4; Mismatches
                                                                                                                                                                                                                                                                                                                                                         Score 23; DB
Pred. No. 89;
                                                                                                                                                                                                                                                                                                                                                                                                                             2 GGTTGCGGAGGGTGGGCCTGGGA 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 GGUUGCGGAGGGUGGGCCUGGGA 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Thompson J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            hTR siNA-target RNA - SEQ ID 92.
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                                                                                                                                                                                                                                                                                                                                                           5.1%;
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2002US-0363124P.
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2002US-0409293P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Beigelman L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         26-FEB-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                          19; Conservative
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                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO2003074654-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unidentified.
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29-AUG-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20-FEB-2002;
11-MAR-2002;
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Example 24; SEQ ID NO 85; 593pp; English.
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                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                       ADG29525;
                                                                                                                                                                                                                                                                                                                                                        RESULT 106
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                 The invention relates to a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene comprising one or more chemical modifications and each strand of the double-stranded siNA comprises about 21 nucleotides. The siNA of the invention demonstrates antiarteriosclerotic, neuroprotective, cootropic, antiparkinsonian and anticonvulsant activities and may be useful for down-regulating the expression of an endogenous mammalian target gene and therefore in the treatment of any disease or condition tissue or organism. The disease or condition any include pulmonary diseases such as restenosis, atherosclerosis, Alzheimer's disease.

Parkinson's disease, epilepsy, dementia, huntington's disease or employed any any include pulmonary disease, epilepsy, dementia, huntington's disease or expensions. The current sequence is that of the siNA target DNA of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New double-stranded short interfering nucleic acid molecule, useful for down-regulating the expression of an endogenous mammalian target gene or for treating diseases that respond to modulation of gene expression or
                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        antiarteriosclerotic; neuroprotective; nootropic; antiparkinsonian; anticonvulsant; pulmonary disease; restenoisis, thetrosclerosis; Alzheimer's; Parkinson's; epilepsy; dementia; huntington's; amyotrophic lateral sclerosis; gene therapy; target; ss; hTR.
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                                                                                                                                                                                                                                           5.1%; Score 23; DB 1; Length 23; 78.3%; Pred. No. 89; 0; Indels iive 5; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              double-stranded short interfering nucleic acid; siNA;
                                                                                                                                                                                                                    Sequence 23 BP; 4 A; 5 C; 9 G; 0 T; 5 U; 0 Other;
Example 24; SEQ ID NO 92; 593pp; English.
                                                                                                                                                                                                                                                                                            298 GCGAAGAGTTGGGCTCTGTCAGC 320
                                                                                                                                                                                                                                                                                                          1 GCGAAGAGUUGGGCUCUGUCAGC 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Thompson J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      hTR siNA-target RNA - SEQ ID 85.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; 2002US-0386782P.
; 2002US-0406784P.
; 2002US-0408378P.
; 2002US-0409293P.
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2002US-0363124P.
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                                                                                                                                                                                                                                                                                                                                                                               ADG29519 standard; RNA; 23
                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                    18; Conservative
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                                                                                                                                                                                                                                                        Local Similarity
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05-SEP-2002;
09-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unidentified
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                                                                                                                                                                                                                                                                                                                                                                                                       ADG29519;
                                                                                                                                                                                                                                              Query Match
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Matches
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The invention relates to a double-stranded short interfering nucleic acid

(siNA) molecule that down-regulates expression of an endogenous mammalian

control gene comprising one or more chemical modifications and each strand

cof the double-stranded siNA comprises about 21 nucleotides. The siNA of

the invention demonstrates antiarteriosclerotic, neuroprotective,

controlot, antiparkinsonian and anticonvulsant activities and may be

cueful for down-regulating the expression of an endogenous mammalian

cueful for down-regulating the expression of an endogenous mammalian

cueful for down-regulating the expression of any disease or condition

trayet gene and therefore in the treatment of any disease or condition

control of the responds to modulation of gene expression or activity in a cell,

clissue or organism. The disease or condition may include pulmonary

diseases such as restenosis, atherosclerosis, Alzheimer's disease,

clissue or organism sections, atherosclerosis, Alzheimer's disease or

samyotrophic lateral sclerosis. Furthermore, the sinA may be utilised for

camput therapy applications. The current sequence is that of the sinA

traget DNA of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1; Length 23;
89;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 23 BP; 2 A; 3 C; 14 G; 0 T; 4 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5.1%; Score 23;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2 GGTTGCGGAGGGTGGGCCTGGGA 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Thompson J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             hTR siNA-target RNA - SEQ ID 91.
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2002US-0363124P.
2002US-0386782P.
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2002US-0408378P.
2002US-0409293P.
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Matches 19; Conservative
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Example 24; SEQ ID NO 87; 593pp; English.
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G Usman N,
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11-MAR-2002;
06-JUN-2002;
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05-SEP-2002;
09-SEP-2002;
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                                                                                                                                                                                                                                                                                                                                                RESULT 108
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                       The invention relates to a double-stranded short interfering nucleic acid (siNN) moblecule that down-regulates expression of an endogenous mammalian target gene comprising one or more chemical modifications and each strand of the double-stranded siNA comprises about 2; nucleotides. The siNA of
                                                                 the invention demonstrates and antiarteriosclerotic, neuroprotective, nootropic, antiparkinsonian and anticonvulsant activities and may be useful for down-regulating the expression of an endogenous mammalian target gene and therefore in the treatment of any disease or condition that responds to modulation of gene expression or activity in a cell, tissue or organism. The disease or condition may include pulmonary diseases such as restenoosis, atherosclerosis, Albrahamer's disease, Parkinson's disease, epilepsy, dementia, huntington's disease or amyotrophic lateral sclerosis. Furthermore, the siNA may be utilised for gene therapy applications. The current sequence is that of the siNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New double-stranded short interfering nucleic acid molecule, useful for down-regulating the expression of an endogenous mammalian target gene or for treating diseases that respond to modulation of gene expression or activity.
                                                                                                                                                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Fosnaugh K;
                                                                                                                                                                                                                                     Score 23; DB 1; Length 23;
Pred. No. 89;
7; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pavco P,
                                                                                                                                                                                                                Sequence 23 BP; 6 A; 7 C; 3 G; 0 T; 7 U; 0 Other;
  Example 24; SEQ ID NO 91; 593pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Beigelman L, Chowrira
                                                                                                                                                                                                                                                                                       146 TTCCACCGTTCATTCTAGAGCAA 168
                                                                                                                                                                                                                                                                                                   Thompson J;
                                                                                                                                                                                                                                                                                                                                                                                                                                              hTR siNA-target RNA - SEQ ID 87.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (SIRN-) SIRNA THERAPEUTICS INC
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2002US-0386782P.
2002US-0406784P.
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2002US-0409293P.
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                                                                                                                                                                                         target DNA of the invention.
                                                                                                                                                                                                                                                                                                                                                                         ADG29521 standard; RNA; 23
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                                                                                                                                                                                                                                      Query Match
Best Local Similarity 69.6
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 J, bers.
c. Usman N,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unidentified
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Jamison S,
                                                                                                                                                                                                                                                                                                                                                                                                ADG29521;
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The invention relates to a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene comprising one or more chemical modifications and each strand of the double-stranded siNA comprises about 21 nucleotides. The siNA of
                                                                                                                               the invention demonstrates antiarteriosclerotic, neuroprotective, nootropic, antiparkinsonian and anticonvulsant activities and may be useful for down-regularing the expression of an endogenous mammalian target gene and therefore in the treatment of any disease or condition that responds to modulation of gene expression or activity in a cell, tissue or organism. The disease or condition may include pulmonary diseases such as restenosis, atherosclerosis, Albreimer's disease, Parkinson's disease, epilepsy, dementia, huntington's disease or amyotrophic lateral sclerosis. Furthermore, the siNA may be utilised for gene therapy applications. The current sequence is that of the siNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1; Length 23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 23 BP; 6 A; 10 C; 6 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       89;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 23;
Pred. No. 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             283 GCACCCACTGCCACCGCGAAGAG 305
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 gcacccacueccacceceaagae 23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           hTR siNA-target RNA - SEQ ID 90.
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2002US-0408378P.
2002US-0409293P.
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2002US-0363124P.
2002US-0386782P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              target DNA of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADG29524 standard; RNA; 23
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Best Local Similarity 95.7
Matches 22; Conservative
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Example 24; SEQ ID NO 86; 593pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                          ADG29522;
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                   The invention relates to a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene comprising one or more chemical modifications and each strand of the double-stranded siNA comprises about 21 nucleotides. The siNA of the invention demonstrates antiarteriosclerctic, neuroprotective, nootropic, antiparkinsonian and anticonvulsant activities and may be useful for down-regulating the expression of an endogenous mammalian that responds to modulation of gene expression or activity in a cell, tissue or organism. The disease or condition way include pulmonary diseases such as restenosis, atherosclerosis, Alzheimer's disease, parkinson's disease, epilepsy, dementia, huntington's disease or amyotrophic lateral sclerosis. Furthermore, the siNA may be utilised for the exapp, applications. The current sequence is that of the siNA
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                                                                                                                                                                                                                                   5.1%; Score 23; DB 1; Length 23; 69.6%; Pred. No. 89;
                                                                                                                                                                                                                                                           0; Indels
                                                                                                                                                                                                            Sequence 23 BP; 4 A; 9 C; 3 G; 0 T; 7 U; 0 Other;
                                                                                                                                                                                                                                                        7; Mismatches
Example 24; SEQ ID NO 90; 593pp; English.
                                                                                                                                                                                                                                                                                   144 CCTTCCACCGTTCATTCTAGAGC 166
                                                                                                                                                                                                                                                                                               1 CCUUCCACCGUUCAUUCUAGAGC 23
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2002US-0406784P.
2002US-0408378P.
2002US-0409293P.
                                                                                                                                                                                                                                                                                                                                                                    ADG29520 standard; RNA; 23 BP.
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2002US-0363124P.
                                                                                                                                                                                      target DNA of the invention.
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                                                                                                                                                                                                                                                           16; Conservative
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                                                                                                                                                                                                                                              Local Similarity
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                                 (siNA) molecule that down-regulates expression of an endogenous mammalian target gene comprising one or more chemical modifications and each strand of the double-stranded siNA comprises about 21 nucleotides. The siNA of
The invention relates to a double-stranded short interfering nucleic acid
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                                                                                                                         the invention demonstrates antiarteriosclerotic, neuroprotective, noctropic, antiparkinsonian and anticonvulsant activities and may be useful for down-regulating the expression of an endogenous mammalian target gene and therefore in the treatment of any disease or condition that responds to modulation of gene expression or activity in a cell, rissue or organism. The disease or condition may include pulmonary diseases such as restenosis, atherosclerosis, Alzheimer's disease. Parkinson's disease, epilepsy, dementia, huntington's disease or amyotrophic lateral sollorosis. Furthermore, the sink may be utilised for gene therapy applications. The current sequence is that of the sink target DNA of the invention.
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89;
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2002US-0406784P.
2002US-0408378P.
2002US-0409293P.
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2002US-0363124P.
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nes 17; Conservative
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gene therapy.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic.
                                                                                                                                                                                                                                                                                                                                                                         AAZ07300;
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                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                        RESULT 112
                                                                                                                                                                                                                              Matches
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                  The invention relates to a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene comprising one or more chemical modifications and each strand of the double-stranded siNA comprises about 21 nucleotides. The siNA of the invention demonstrates antiatretriosclerotic, neuroprotective, casful for down-regulating the expression of an endogenous mammalian that responds to modulation of gene expression of any disease or condition that responds to modulation of gene expression or activity in a cell, cissue or organism. The disease or condition may include pulmonary diseases such as restenosis, atherosclerosis, Alzheimer's disease, platepsy, dementia, huntington's disease or comparations of disease, epilepsy, dementia, huntington's disease or comparations. The current sequence is that of the sinA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This invention relates to a novel method concerning prognostic markers associated with EGFR (epidermal growth factor receptor) positive cancer. Specifically, it refers to a gene expression profiling method that can provide a prediction as to whether a patient is likely to respond well to treatment with an EGFR inhibitor. The present invention describes the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Prognosing a patient with EGFR-expressing colon cancer comprises subjecting a sample comprising EGFR-expressing cancer cells to quantitative analysis of the expression level of the RNA transcript of at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PCR primer to amplify a human cancer prognostic marker DNA SeqID 345.
                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  epidermal growth factor receptor; cancer; gene expression profiling;
microarray; head and neck cancer; colon cancer; metastatic spread;
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                                                                                                                                                                                                                                                          5.1%; Score 23; DB 1; Length 23;
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                                                                                                                                                                                                                                  Sequence 23 BP; 2 A; 8 C; 9 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        human; primer; PCR; prognostic marker; EGFR;
                                                                                                                                                                                                                                                                      Pred. No. 89;
4; Mismatches
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 Example 24; SEQ ID NO 88; 593pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 54; SEQ ID NO 345; 113pp; English.
                                                                                                                                                                                                                                                                                                             395 GCGCGCCGATTCCCTGAGCTG 417
                                                                                                                                                                                                                                                                                                                            GCGCGCGCGAUCCCUGAGCUG 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (VALL-) VALL HEBRON UNIV HOSPITAL
                                                                                                                                                                                                                                                                                                                                                                                                     ADP27908 standard; DNA; 23 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15-NOV-2002; 2002US-0427090P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    least one gene e.g., CD44v3.
                                                                                                                                                                                                         target DNA of the invention.
                                                                                                                                                                                                                                                                      82.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (GENO-) GENOMIC HEALTH INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                      19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              neoplastic disease; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cronin MT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2004-420643/39.
                                                                                                                                                                                                                                                                         Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO2004046386-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                        26-AUG-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           03-JUN-2004.
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                                                                                                                                                                                                                                                                                                                                                                                                                              ADP27908;
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quantitative analysis of the expression level of the RNA transcript of at least one gene selected from the group of CD44v3, CD44v6, DR5, GRO1, KRT17, LAMC2 or their products thereof. It further provides a CDNA microarray containing named genes that represent prognostic transcripts which are useful for determining whether a patient diagnosed with an EGFR expressing head or neck cancer or colon cancer exhibits elevated or decreased expression levels of these genes compared to normal. As such, these methods are also useful for prognosing or predicting the likelihood of cancer-attributable death or progression, including recurrence and metastatic spread of a neoplastic disease, as well as drug resistance. This oligonuclectide sequence is a PCR primer used to amplify a human PCR amplicon DNA sequence used as a prognostic cancer marker, given in an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mouse and human telomerase RNA gene promoters, useful for tumor specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Telomerase RNA; TR; promoter; cytotoxin; cancer; neoplasia; hTR; gene therapy; thymidine kinase gene; anticancer therapy; human; mutagenesis; PCR primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1; Length 23;
89;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 23 BP; 6 A; 9 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 23;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           404 GATTCCCTGAGCTGTGGGACGTG 426
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        23 GATTCCCTGAGCTGTGGGACGTG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  exemplification of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     22-OCT-1999 (first entry)
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Les 23; Conservative
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Sequence 27 BP; 2 A; 4 C; 12 G; 9 T; 0 U; 0 Other;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RNA components of (non)human mammalian telomerase(s) - useful in studying cell senescence and immortalisation.
(especially pGT62-codAupp) comprising the promoter and a viral thymidine kinase gene can be used to convert a prodrug, e.g. gancyclovir, so that neoplasia can be used to crareated. Direct down-regulation of telomerase RNA gene through manipulation of transcription factors may be effective anticancer therapy and the cloning of the hTR gene promoter allows the analysis of therapeutic molecules which modulate hTR promoter activity. Sequences AAZO7696-321 represent PCR primers used in cloning and mutagenisis of human TR gene (hTR) promoter region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Telomerase; mammal; antisense; triplex forming oligonucleotide; plasmid;
                                                                                                                                                                                                                              Gaps
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                                                                                                                                                                                         5.1%; Score 23; DB 1; Length 25;
100.0%; Pred. No. 1e+02;
ive 0; Mismatches 0; Indels
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                                                                                                                                                           Sequence 25 BP; 1 A; 5 C; 15 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Primer used for producing telomerase probe.
                                                                                                                                                                                                                                                                                    GGGTTGCGGAGGGTGGGCCTGGG 25
                                                                                                                                                                                                                                                              1 GGGTTGCGGAGGGTGGGCCTGGG 23
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B;
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95US-00387524.
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                                                                                                                                                                                                                            23; Conservative
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                                                                                                                                                                                                            Best Local Similarity
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GERON CORP
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13-FEB-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
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                                                                                                                                                                                                                                                                                                               Telomerase; mammal; antisense; triplex forming oligonucleotide; plasmid;
                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Marhuenda MA;
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                                                                                                                                                                                                                                                                                Antisense oligonucleotide (P3) inhibiting telomerase activity.
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              Length 27;
                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Andrews WH, Avillon AA, Feng J, Funk W, Greider C,
Villeponteau B;
DB 1; Leus
1.1e+02;
0;
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           Query Match 5.1%; Score 23; DB Best Local Similarity 100.0%; Pred. No. 1.1 Matches 23; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cell senescence and immortalisation.
                                                                            44
                                                                                                       GGAGGGGTGGTGCCATTTTTG 27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       46 CTAACCCTAACTGAGAAGGGCG 67
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95US-00387524.
95US-00485778.
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                                                                                                                                                                                                                                                                                                                                 probe; primer; ribozyme; ss.
                                                                                                                                                                                     AAT11033 standard; DNA; 22
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07-JUN-1995;
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                                                                                                                                                                                                                                                                                                                                                               Synthetic.
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                                                                                                                                                      RESULT 114
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RNA component of mammalian telomerase, esp. human - useful in identifying
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RNA component of mammalian telomerase antisense oligonucleotide P3.
         mammalian, telomerase, antisense oligonucleotide; inhibition; neoplastic; cells; activity; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RNA component; mammalian; telomerase; antisense oligonucleotide;
triple helix; inhibition; neoplastic; cells; activity; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 22; DB 1; Length 22;
Pred. No. 1e+02;
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                                                                                                                                                                                                                                       Andrews WH;
                                                                                                                                                                                                                                                                                                      e.g. candidate telomerase-modulating agents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                              Disclosure; Page 38; 114pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22 AACTGAGAAGGGCGTAGGCGCC 1
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94US-00330123.
95US-00472802.
95US-00482115.
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94US-00330123.
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95US-00482115.
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                                                                                                                                                                                                                (GERO-) GERON CORP.
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Matches 22; Conserv
                                                                                                                                                                                                                                                                                                                                                                                             neoplastic cells
           RNA component;
triple helix;
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                                                                                                                         06-JUL-1995;
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07-JUN-1995;
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                                                 Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The RNA components of (non) human mammalian telomerase(s) especially from mouse, rat and chinese hamster are all claimed. Antisense obligonucleotides can be used to block the activity of the telomerase; probes and primers can be used in detection; vectors and host cells transformed with the isolated telomerase genes can be used for production of telomerases; RNA and DNA ribozymes and triplex forming oligonucleotides directed against the telomerase genes can be used therapeutically as can plasmids. A mouse which lacks the telomerase gene therapeutically as can plasmids. A mouse which lacks the telomerase gene the role it plays in immortalisation. The antisense oligonucleotide is synthesised as a 2-0-methyl RNA oligonucleotide and is more resistant to hydrolysis than unmodified RNA oligonucleotides (See AATI1032-35)
                                                                                                                                                                                                                                                                                                                                                                                                                                                components of (non)human mammalian telomerase(s) - useful in studying
                                                                                                                                     Telomerase; mammal; antisense; triplex forming oligonucleotide; plasmid;
                                                                                                                                                                                                                                                                                                                                                                                 Marhuenda MA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RNA component of mammalian telomerase antisense oligonucleotide TA3.
                                                                                                             Antisense oligonucleotide (TA3) inhibiting telomerase activity.
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100.0%; Pred. No. 1e+02;
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                                                                                                                                                                                                                                                                                                                                                                                 Greider C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 22 BP; 2 A; 9 C; 5 G; 0 T; 6 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.08; Prea. ...
                                                                                                                                                                                                                                                                                                                                                                                 Funk W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                              cell senescence and immortalisation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 23; 85pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        54 AACTGAGAAGGGCGTAGGCGCC 75
                                                                                                                                                                                                                                                                                                                                                                                   Feng J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AACTGAGAAGGGCGTAGGCGCC 1
                                                                                                                                                                                                                                                                                                                                            COLD SPRING HARBOR LAB
                                                                                                                                                                                                                                                                             94US-00272102.
94US-00330123.
95US-00387524.
95US-00485778.
                                     AAT11034 standard; DNA; 22 BP.
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                                                                                                                                                                                                                                                      95WO-US008620
                                                                                                                                                    probe; primer; ribozyme; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAT10288 standard; DNA; 22
                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                         Villeponteau B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1996-097428/10.
                                                                                                                                                                                                                                                                                                                                                         GERON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                    WO9601614-A2
                                                                                                                                                                                                                                                      07-JUL-1995;
                                                                                                                                                                                                                                                                                          27-OCT-1994;
13-FEB-1995;
                                                                                      02-JUL-1996
                                                                                                                                                                                                                                                                                                                    07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 09-SEP-1996
                                                                                                                                                                                                                             25-JAN-1996
                                                                                                                                                                             Synthetic.
                                                           AAT11034
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                                                                                                                                                                                                                                                                                                                                            (COLD-)
                                                                                                                                                                                                                                                                                                                                                         (GERO-)
                      Matches
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AAT10288/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SXXXXX
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Query Match
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                                                                              Best Loc
Matches
                                                                                                                                                                   RESULT 119
                                                                                                                                                                             AAT58812/
SSSSS
                                                                                                                                                                                                             ð
                                                                                                                              원
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                               RNA component of mammalian telomerase, esp. human - useful in identifying e.g. candidate telomerase-modulating agents.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RNA component of mammalian telomerase, esp. human - useful in identifying
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present sequence, a return primer for the RNA component of human telomerase (RCHT), was used in a primed in situ labelling (RRINS) procedure. The RCHT can be used in the recombinant prodn of an active telomerase mol., capable of adding sequences to chromosomal DNA telomerase and in the synthesis of mutant sequences for the detection of mutant mammalian telomerase RNA component polynucleotides. The RCHT may also be used in the identification of telomerase modulating agents, and in the detection of telomerase related, or neoplastic conditions in a patient. Polynucleotides of at least 25 consecutive nucleotides
                                                                                      The present sequence is a RNA component of mammalian telomerase, antisense oligonucleotide, which can be used, along with triple helix forming sequences, to inhibit telomerase activity in cells, esp. neoplastic cells
                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                            recombinant production, synthesis, mutant, detection, mammalian, identification, modulating agent, neoplastic condition; transcriptional regulatory sequence, gene therapy, disease;
                                                                                                                                                                                        ö
                                                                                                                                                                 4.9%; Score 22; DB 1; Length 22; 100.0%; Pred. No. 1e+02; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                   component; human; telomerase; return primer; PRINS;
                                                                                                                                                                                                                                                                                                                                                           RNA component of human telomerase PRINS return primer.
                                                                                                                                             Sequence 22 BP; 3 A; 6 C; 6 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Andrews WH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                candidate telomerase-modulating agents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 13; Page 90; 114pp; English.
                                                                Disclosure; Page 38; 114pp; English
                                                                                                                                                                                                             46 CTAACCCTAACTGAGAAGGGCG 67
                                                                                                                                                                                                                          CTAACCCTAACTGAGAAGGGCG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Funk W,
                                                                                                                                                                                                                                                                               AAT10308/c
ID AAT10308 standard; DNA; 22 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     94US-00330123.
95US-00472802.
95US-00482115.
                                                                                                                                                                                                                                                                                                                                                                                                                               primed in situ labelling; ss.
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                                                                                                                                                                                       22; Conservative
         WPI; 1996-097581/10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1996-097581/10.
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                                                                                                                                                                            Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      06-JUL-1995;
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07-JUN-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic
                                                                                                                                                                                                                                  22
                                                                                                                                                                                                                                                                                                                AAT10308;
                                                                                                                                                                  Query Match
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                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                   RNA
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The present sequence represents PCR 3'-primer hal88 used for amplifying
the human telomerase (hTR). The RNA and DNA can be used in hybridisation
cassays to detect or quantify telomerase activity in cells, tissue or
fluid samples, e.g. for diagnosis of eukaryotic parasites (yeast and
protozoa) or tumours. It is also useful as primers for amplification
cassays. The truncated or recombinant vertebrate telomerase is used
therapeutically to increase telomerase activity (also as reagents in the
screening assay) while the RNA or other inhibitors such as antisense
molecules, are used to reduce such activity. Typical applications are
initiation/restoration of activity to cause senescence or to prevent
immortalisation of cells in tumours or parasites. The DNA is also used to
produce recombinant telomerase, which can then be used conventionally to
craise antibodies for diagnostic detection of telomerase. Detecting
call cause fewer side effects than drugs currently used to treat such
infections. The DNA encodes those parts of the RNA essential for activity
his are significantly changer than argus currently used to treat such
the but and a parasite of the manning and activity of the parts of the
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identical, or complementary to the RCHT sequence linked to heterologous transcriptional regulatory sequences, can be used for the gene therapy of human diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cancer; eukaryotic parasite; hTR; vertebrate telomerase; yeast; protozoa; tumour; antibody; polymerase chain reaction; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA encoding essential RNA components of human telomerase - also truncated or recombinant telomerase, useful for diagnosis and treatment of cancer and infection by eukaryotic parasites.
                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    out are significantly shorter than the endogenous RNA component
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                                                                                                                                                                                                                                                                                                      4.9%; Score 22; DB 1; Length 22;
100.0%; Pred. No. 1e+02;
ive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                Sequence 22 BP; 5 A; 5 C; 12 G; 0 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human telomerase PCR 3'-primer hal88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       183 CTGCTGGCCCGTTCGCCCCTCC 204
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CIGCIGGCCGTTCGCCCCTCC 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Autexier C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1997-099928/09
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nes 22; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic.
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Synthetic
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                                                                                                     AAZ23628;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present sequence represents an antisense oligonucleotide directed against the human telomerase RNA component gene sequences. The oligonucleotide can be used as an affinity agent in the methods of the invention, which are used to purify human telomerase. The methods of the use of several sequential steps, including the use of two matrixes that bind molecules bearing negative charges, a matrix that binds molecules bearing positive charges, an affinity purification step and a size separation. Telomerase is a particular target of anticancer therapies, and is useful in assays for characterizing (pre) cancerous cells. Telomerase can also be used to screen for specific modulators, for biochemical analysis of its activity, and in preparation of antibodies. Fragments of telomerase, or nucleic acid encoding them, are used in vaccines, and for treating over expression of telomerase, particularly in
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                                                                                                                                                                                                                                                                                                                                                            Human; telomerase RNA component; anticancer therapy; purification; assay; vaccine; cancer; antisense oligonuclectide; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pruzan RA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Purification of telomerase on affinity material - useful for, e.g. diagnosis and treatment of cancer.
                                                                                                                                                                                                                                                                                                                            Antisense oligonucleotide P3 for human telomerase RNA component.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .
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Pred. No. 1e+02;
0; Mismatches 0; Indels
  Indels
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100.0%; Pred. No. 10.
0, Mismatches
  Mismatches
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/note= "biotinylated"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                                            163 GAGCAAACAAAAATGTCAGCT 184
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure, Page 24; 76pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             46 CTAACCCTAACTGAGAAGGGCG 67
                                                                          GAGCAAACAAAAATGTCAGCT
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                                                                                                                                                                                            AAV63646 standard; DNA; 22
                                                                                                                                                                                                                                                                                  (first entry)
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  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 22; Conserve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1998-594485/50
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Key
modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             04-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     04-APR-1997;
                                                                                                                                                                                                                                                                                    15-FEB-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO9845450-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15-OCT-1998.
    22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kealey JT;
                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic.
                                                                                                                                                                                                                                         AAV63646;
                                                                                     22
      Matches
                                                                                                                                                                        AAV63646/A

AAV65AAV6

AAV6

AAV6

XX

AAV6

XX

AAV6

XX

BDB

ADD

BDB

CCC

CC
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This invention describes a novel composition comprising human telomerase having at least 2000-fold (preferably at least 6000-fold) increased treative purity compared with crude extract of cells from adenovirustransformed kidney cell line. The composition is useful for eliciting an immune response in animals and may therefore be used as a vaccine for treating diseases associated with the overexpression of telomerase e.g. cancer. AAZ23656-Z23637 represent oligonucleotides used in the isolation of human clone 28-1 which contains a fragment of the human telomerase
                                                                                                                                                                               human; immune response; cancer; vaccine; treatment; disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Compositions comprising human telomerase, useful for treating diseases associated with overexpression of telomerase e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Kealey JT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         .
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Antisense oligonucleotide for human telomerase, BIOTIN P3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4.9%; Score 22; DB 1; Length 22;
100.0%; Pred. No. 1e+02;
lve 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pruzan RA,
                                                                                                                                   Human clone 28-1 telomerase oligonucleotide oligo-P3.
                                                                                                                                                                                                                                                                                                                                                                          /*tag= a
/note= "5'-biotinylated cytidine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 22 BP; 3 A; 6 C; 6 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Weinrich SL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      described in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Col 43-44; 34pp; English.
                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4.>°,
100.0%; Pre-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          46 CTAACCCTAACTGAGAAGGGCG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Lichtsteiner SP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP.
AAZ23628 standard; DNA; 22 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      97US-00833377.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     95US-00510736.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAS09473 standard; DNA; 22
                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity 100.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1999-590379/50.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORP.
                                                                                                                                                                                                                                                                                                                               Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (GERO-) GERON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         04-APR-1997;
                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       04-AUG-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Atkinson EM,
Vasserot AP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                24-OCT-2001
                                                                                          07-JAN-2000
                                                                                                                                                                                      Telomerase;
                                                                                                                                                                                                                                                                                                                                                                                                                                               US5968506-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19-OCT-1999
                                                                                                                                                                                                                 primer; ss.
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XEXEXEX
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22 CTAACCCTAACTGAGAAGGGCG 1

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(UYRP ) UNIV ROCHESTER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Telomerase PCR primer.
                                                                                                                                                                                                                                         WPI; 2003-403289/38.
                                                                     WO2003034985-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO2003034985-A2
                                                Ното варіеля
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                             01-MAY-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-MAY-2003
                                                                                                                                                                                                                  Ч,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ACC57543;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                 Rowley
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ò
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                           The sequence represents a biotinylated antisense oligonucleotide used in the purification of human telomerase. The invention relates to a purified buman telomerase core enzyme protein comprising 2000-fold increased purity compared with a crude extract of cells from adenovirus-transformed kidney cell line (293 cells) and when associated with telomerase RNA component has DNA polymerase activity and a molecular weight of 200-2000 kilo Daltons (kDa). The purified telomerase is useful for inducing a humoral or cell-mediated immune response in an animal. Purified telomerase or immunogenic fragments are useful as vaccines for treating diseases associated with over-expression of telomerase, which are useful as affinity agents in isolating the proteins and for detecting the presence of proteins in a sample, such as cellor tissue. Identification of planerase is detecting the presence of proteins in a sample, such as cellor tissue. Identification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      of telomerase aids in diagnosis of cancer or pre-cancerous states. Telomerase and/or telomerase associated proteins are also useful for screening compounds to identify agents that alter the association of telomerase-associated proteins, such as nucleolin or EF2H with telomerase
                                                                                                                                                                                                                                                                                                                          Purified human telomerase, useful for inducing immune response in animals, comprises several thousand folds increased purity compared with
                                                                                                                                                                                                                                                                 Pruzan RA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Telomerase; enzyme; RNA interference; short interfering RNA; siRNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
Human, Telomerase, vaccine, antibody, cancer; EF2H, nucleolin, antisense oligonucleotide, BIOTIN P3; ss.
                                                                                                                                                                                                                                                                 Vasserot AP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 22; DB 1; Length 22;
Pred. No. 1e+02;
0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 22 BP; 3 A; 6 C; 6 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                 Atkinson EM, Lichtsteiner SP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4.9%; Score ...
100.0%; Pred. No. 1e+0;
                                                                                 /*tag= a
/mod_base= C
/note= "C is biotinylated"
                                                                                                                                                                                                                                                                                                                                                   cytoplasmic crude cell preparations.
                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          46 CTAACCCTAACTGAGAAGGGCG 67
                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Col 18; 29pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22 craacccraacreagaagggg 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ACC57544 standard; DNA; 22 BP
                                                                                                                                                                                99US-00420056
                                                                                                                                                                                                     95US-00510736.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 100.
Matches 22; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Telomerase PCR primer.
                                                                                                                                                                                                                                                                                                    WPI; 2001-450477/48.
                                                                                                                                                                                                                                         (GERO-) GERON CORP.
                                                          Key
modified_base
                                                                                                                                                                                                                                                             Weinrich SL,
                                     Homo sapiens
                                                                                                                                JS6261556-B1
                                                                                                                                                                                18-OCT-1999;
                                                                                                                                                                                                      04-AUG-1995;
04-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28-JUL-2003
                                                                                                                                                       17-JUL-2001.
                                                                                                                                                                                                                                                                            Kealey JT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ACC57544;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 123
 8
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The present sequence is a PCR primer used for RT-PCR quantitation of telomerase RNA (see also ACC5751). The invention relates to the discovery that double-stranded interfering RNAs, such as short interfering RNAs (SIRNA), which target telomerase RNA or telomerase reverse transcriptase (TERT) mRNA, are capable of inhibiting telomerase activity. In cancer cells, inhibition of telomerase leads to telomere shortening, end-to-end chromosomal fusion, and apoptosis Telomerase inhibition can also be used for treatment of infertility, for contraception or sterilisation, for immunosuppression, for treatment of yeast, parasite and fungal infections, and in antiinflammatory therapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel nucleic acid encoding or comprising interfering RNAs which target telomerase RNA, useful for inhibiting telomerase activity for treating cancer, infertility and disorders of the immune system.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
cancer; tumour; cytostatic; contraceptive; immunosuppressive; antiinfertility; fungicide; antiparasitic; antiinflammatory; human; gene therapy; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Telomerase; enzyme; RNA interference; short interfering RNA; siRNA; cancer; tumour; cytostatic; contraceptive; immunosuppressive; antiinfertility; fungicide; antiparasitic; antiinflammatory; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
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100.0%; Pred. No. 1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 22 BP; 6 A; 7 C; 7 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                176 ATGTCAGCTGCTGGCCCGTTCG 197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             22 ATGTCAGCTGCTGGCCCGTTCG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 2; Page 27; 52pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ACC57543 standard; DNA; 22 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    gene therapy; PCR; primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16-OCT-2002; 2002WO-US033146.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20-FEB-2002; 2002US-0359196P. 22-MAY-2002; 2002US-0383195P.
                                                                                                                                                                                                                                                                                                                                                                                                     16-OCT-2002; 2002WO-US033146.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22-OCT-2001; 2001US-0345326P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.08;
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Les 22; Conservative
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This invention relates to a purified human telomerase protein, which when associated with telomerase RNA component has DNA polymerase activity.

Also disclosed in the specification is a method for assessing a regulator (preferably a telomerase inhibitor or activator of telomerase involves measuring telomerase enzymatic activity of the composition in presence of a regulator. The telomerase protein of the invention may be used in the developing and testing assays for measuring telomerase activity which are useful in characterising cancer and pre-cancer cells, for identifying and testing aspainst telomerase. The mammalian telomerase protein of the invention is at least approximately 3000 fold more pure (in terms of telomerase activity per weight of protein) than a crude extract of the invention is at least approximately 3000 fold more pure (in terms of telomerase activity per weight of protein) than a crude extract of cell from adenovirus- transformed kidney cell. Purified telomerase for facilitates a thorough biochemical analysis of the enzyme's mechanism for developing mechanism-based regulators. The present sequence represents a human telomerase antisense oligonucleotide which has affinity to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Obtaining telomerase, by preparing enriched solution from cell expressing telomerase, combining the solution with oligonucleotide having specific affinity for the protein and collecting protein bound to oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                        telomerase and is used to purify the telomerase protein of the invention
Composition useful e.g. in diagnosis of cancer comprises complex of telomerase protein with telomerase RNA component.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 mammalian telomerase protein; telomerase purification; telomere; anion exchange matrix; cation exchange matrix; selectivity matrix; gel filtration chromatography; gradient centrifugation; antisense oligonucleotide; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human telomerase RNA component antisense oligonucleotide seq id 2.
                                                                                                                                                                                                                                                                                                                                                                                                                        4.9%; Score 22; DB 1; Length 22;
100.0%; Pred. No. 1e+02;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                            Sequence 22 BP; 3 A; 6 C; 6 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Lichtsteiner SP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; SEQ ID NO 2; 24pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 46 CTAACCCTAACTGAGAAGGGCG 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22 craacccraacreagaagggg 1
                                                   English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            95US-00510736.
97US-00833377.
99US-00420056.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ВР.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20-NOV-2000; 2000US-00717829.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADC35649 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 4.9
Best Local Similarity 100.
Matches 22; Conservative
                                                       Disclosure; Col 9; 24pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Atkinson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2003-742824/70.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (GERO-) GERON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Weinrich SL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US6545133-B1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              04-AUG-1995;
04-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       L8-DEC-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADC35649;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADC35649,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ð
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
                                                                                                                                                                                                                                                                        The present sequence is a PCR primer used for RT-PCR quantitation of telomerase RNA (see also ACC57551). The invention relates to the discovery that double-stranded interfering RNAs, such as short interfering RNAs (sIRNA), which target telomerase RNA or telomerase reverse transcriptase (fIRNA), and capable of inhibiting telomerase activity. In cancer cells, inhibition of telomerase leads to telomera shortening, end-to-end chromosomal fusion, and apoptosis. Telomerase inhibition can also be used for treatment of infertility, for contraception or sterilisation, for immunosuppression, for treatment of yeast, parasite and fungal infections, and in antiinflammatory therapies
                                                                                                                                                                        Novel nucleic acid encoding or comprising interfering RNAs which target telomerase RNA, useful for inhibiting telomerase activity for treating cancer, infertility and disorders of the immune system.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pruzan RA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Vasserot AP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4.9%; Score 22; DB 1; Length 22;
.00.0%; Pred. No. 1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human telomerase antisense oligonucleotide primer P3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 22 BP; 2 A; 3 C; 11 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Telomerase; antisense; primer; P3; ss; cancer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /*tag= a
/note= "Biotinylated"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CTGGGAGGGGTGGTGGCCATTT 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CTGGGAGGGGTGGTGGCCATTT 22
                                                                                                                                                                                                                                               Example 2; Page 27; 52pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     95US-00510736.
97US-00833377.
99US-00420056.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.08;
       22-OCT-2001; 2001US-0345326P.
20-FEB-2002; 2002US-0359196P.
22-MAY-2002; 2002US-0383195P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABX10983 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        22; Conservative
                                                                            (UYRP ) UNIV ROCHESTER
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                                                                                                                                            WPI; 2003-403289/38
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Weinrich SL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      04-AUG-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     JS6517834-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        04-APR-1997;
18-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17-AUG-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     11-FEB-2003
                                                                                                            Rowley PT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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ABX10983/
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Gaps

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RA;

Pruzan

Vasserot AP,

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The invention describes a method of obtaining mammalian telomerase protein (1). The method involves preparing enriched solution (BS) from a cell expressing telomerase where the component of (1) in ES is separated from other proteins expressed by cell by combining ES with coligonucleotide (0) having specific affinity for (1), and collecting protein bound to (0). The oligonucleotide comprises a retrievable label such as biotin and contains a sequence that is specifically recognized by telomerase protein The oligonucleotide contains or does not contain the sequence (TTAGGG)3. The method further comprises combining a fraction containing telomerase protein with a maion exchange matrix, and collecting protein with a cation exchange matrix. (such as a heparin matrix), and collecting protein that binds the matrix (such as a heparin comprises successively enriching fractions containing telomerase protein that binds the matrix. The method on several different ion exchange matrices and combining a fraction containing telomerase protein with an intermediate selectivity matrix.

Collecting protein that binds the matrix, where the intermediate selectivity matrix, collecting protein that binds the matrix, where the intermediate selectivity matrix, and separating a fraction containing the telomerase protein by gel filtration chromatography or gradient centrifugation. The relomerase is enriched from an extract of cells stably expressing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
                                                                                                                                                                                                                                                                                                                                               telomerase. This sequence represents an antisense oligonucleotide to the
                                                                                                                                                                                                                                                                                                                                                              component of human telomerase that can be used in the purification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pruzan RA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Telomerase activity; therapy; cancer; cytostatic; antisense; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                               4.9%; Score 22; DB 1; Length 22;
100.0%; Pred. No. 1e+02;
ve 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human telomerase RNA antisense oligonucleotide, P3.
                                                                                                                                                                                                                                                                                                                                                                                                               Seguence 22 BP; 3 A; 6 C; 6 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note= "Biotin labelled"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  46 CTAACCCTAACTGAGAAGGCG 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CTAACCCTAACTGAGAAGGGCG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           J= a
base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 Similarity 100.0%; Pr
22; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADG62871 standard; DNA; 22 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              04-AUG-1995; 95US-00510736.
04-APR-1997; 97US-0083377.
18-OCT-1999; 99US-00420056.
20-NOV-2000; 2000US-00717828.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               24-DEC-2002; 2002US-00330872
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Weinrich SL, Atkinson EM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (WEIN/) WEINRICH S L.
(ATKI/) ATKINSON E M.
(LICH/) LICHTSTEINER S P.
(VASS/) VASSEROT A P.
(PRUZ/) PRUZAN R A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DOE/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
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                                                                                                                    The invention relates to a method for identifying regulators of telomerase activity that may be useful for treating cancers. The method may be used to identify regulators e.g. antibodies, of telomerase activity which may be useful as cancer treatments. It has been found that found that the cells of many human cancers have telomerase activity. This helps explain why cancer cells continue dividing without becoming senescent. If telomerase activity in cancer cells can be inhibited, the cancer cells are expected to reach senescence and cease dividing. The present sequence is human telomerase antisense oligonulceotide used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Telomerase; enzyme; RNA interference; short interfering RNA; siRNA; telomerase; cancer; tumour; cytostatic; contraceptive; immunosuppressive; antiinfertility; fungicide; antiparasitic; antiinflammatory; human; gene therapy; PCR; primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            nucleic acids which are complementary to each other and to target nucleic acid e.g., telomerase RNA or mRNA encoding telomerase reverse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel double-stranded short interfering RNA having sense and antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                       Gapa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present sequence is a PCR primer used for RT-PCR quantitation of telomerase RNA (see also ACC58040). The invention relates to the discovery that double-stranded interfering RNAs, such as short interfering RNAs (sIRNA), which target telomerase RNA or telomerase
                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                        Identifying telomerase regulators useful for treating cancer.
                                                                                                                                                                                                                                                                                                                                                                                          4.9%; Score 22; DB 1; Length 22;
100.0%; Pred. No. 1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Indels
                                                                                                                                                                                                                                                                                                                                                 Sequence 22 BP; 3 A; 6 C; 6 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Mismatches
                                                                                                                                                                                                                                                                                                            illustrate the method of the invention.
                                                                               Disclosure; SEQ ID NO 2; 22pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              46 CTAACCCTAACTGAGAAGGGCG 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 2; Page 24; 37pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CTAACCCTAACTGAGAAGGGCG 1
                                                                                                                                                                                                                                                                                                                                                                                                               100.08;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22-OCT-2001; 2001US-0345326P.
20-FEB-2002; 2002US-0359196P.
22-MAY-2002; 2002US-0383195P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ACC58032 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Telomerase PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (UYRP ) UNIV ROCHESTER
WPI; 2003-811733/76.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2003-403336/38.
                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
nes 22; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO2003035667-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ACC58032;
                                                                                                                                                                                                                                                                                                                                                                                              Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 128
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reverse transcriptase (TERT) mRNA, are capable of inhibiting telomerase activity. In cancer cells, inhibition of telomerase leads to telomere shortening, end-to-end chromosomal fusion, and apoptosis. Telomerase inhibition can also be used for treatment of infertility, for contraception or sterilisation, for immunosuppression, for treatment of yeast, parasite and fungal infections, and in antiinflammatory therapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    telomerase; cancer; tumour; cytostatic; contraceptive; immunosuppressive; antiinfertility; fungicide; antiparasitic; antiinflammatory; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel double-stranded short interfering RNA having sense and antisense unductaic acids which are complementary to each other and to target nucleic acid e.g., telomerase RNA or mRNA encoding telomerase reverse transcriptase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     contraception or sterilisation, for immunosuppression, for treatment of yeast, parasite and fungal infections, and in antiinflammatory therapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present sequence is a PCR primer used for RT-PCR quantitation of telomerase RNA (see also ACC58040). The invention relates to the discovery that double-stranded interfering RNAs, such as short interfering RNAs, such as short interfering RNAs (SIRNA), which target telomerase RNA or telomerase reverse transcriptase (TERT) mRNA, are capable of inhibiting telomerase activity. In cancer cells, inhibition of telomerase leads to telomera shortening, end-to-end chromosomal fusion, and apoptosis. Telomerase inhibition can also be used for treatment of infertility, for contraception or sterilisation, for immunosuppression, for treatment of
                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Telomerase, enzyme; RNA interference; short interfering RNA; siRNA;
                                                                                                                                                                                                                          ;
                                                                                                                                                                                 DB 1; Length 22;
1e+02;
                                                                                                                                                                                                                        0; Indels
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                                                                                                                                       Sequence 22 BP; 6 A; 7 C; 7 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                             Query Match
4.9%; Score 22; DB
Best Local Similarity 100.0%; Pred. No. 1e+
Matches 22; Conservative 0; Mismatches
                                                                                                                                                                                                                                                        176 ATGTCAGCTGCTGGCCCGTTCG 197
                                                                                                                                                                                                                                                                                               22 ArgrcAgcrecreeccerres 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 2; Page 24; 37pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            gene therapy; PCR; primer; ss.
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2002US-0359196P.
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                                                                                                                                                                                                                                                                                                                                                                                            ACC58031 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (UYRP ) UNIV ROCHESTER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Telomerase PCR primer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        mouse, rat and chinese master are all claimed. Antisense oligonucleotides can be used to block the activity of the telomerase; probes and primers can be used in detection; vectors and host cells transformed with the isolated telomerase genes can be used for production of telomerases; RNA and DNA ribozymes and triplex forming of telomerases; RNA and DNA ribozymes and triplex forming oligonucleotides directed against the telomerase genes can be used therapeutically as can plasmids. A mouse which lacks the telomerase gene (also claimed) can be used for study of telomere regulation in vivo, and AATI1059, AATI1060 which are complementary to human telomerase RNA component sequences can be used to identify and amplify homologous sequences from other non-human mammals. The amplified fragments can then be used as probes to identify telomerase genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RNA components of (non)human mammalian telomerase(s) - useful in studying cell senescence and immortalisation.
                                                                                                                                                                                                                                     Telomerase; mammal; antisense; triplex forming oligonucleotide; plasmid;
     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Marhuenda MA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The RNA components of (non) human mammalian telomerase(s) especially
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0
    Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Greider C,
                                                                                                                                                                                                          Primer used for amplifying telomerase RNA fragments.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 21 BP; 5 A; 5 C; 11 G; 0 T; 0 U; 0 Other;
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 Mismatches
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                                                      22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Feng J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          184 TGCTGGCCCGTTCGCCCTCC 204
                                                       CTGGGAGGGGTGGTGGCCATTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TGCTGGCCCGTTCGCCCCTCC 1
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0
                           19 CTGGGAGGGGTGGTGGCCATTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                              (COLD-) COLD SPRING HARBOR LAB
                                                                                                                             H.
                                                                                                                                                                                                                                                                                                                                                                                         94US-00272102.
94US-00330123.
95US-00387524.
95US-00485778.
                                                                                                                                                                                                                                                                                                                                                                 95WO-US008620
                                                                                                                                                                                                                                                    probe; primer; ribozyme; ss.
                                                                                                               .058/c
AAT11058 standard; DNA; 21
                                                                                                                                                                                 (first entry)
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Avillon AA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1996-097428/10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (GERO-) GERON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
21; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Villeponteau B;
                                                                                                                                                                                                                                                                                                         WO9601614-A2
                                                                                                                                                                               02-JUL-1996
                                                                                                                                                                                                                                                                                                                                                                 07-JUL-1995;
                                                                                                                                                                                                                                                                                                                                                                                         07-JUL-1994;
27-OCT-1994;
 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                  12-VUL-1995;
                                                                                                                                                                                                                                                                                                                                     25-JAN-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Andrews WH,
                                                                                                                                                                                                                                                                                Synthetic.
                                                                                                                                                     AAT11058;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21
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 Matches
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Matches
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4.9%; Score 22; DB 1; Length 22; 100.0%; Pred. No. 1e+02;

Query Match Best Local Similarity 96WO-US009517. 95US-00478352.

Autexier C;

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DNA encoding essential RNA components of human telomerase - also truncated or recombinant telomerase, useful for diagnosis and treatment of cancer and infection by eukaryotic parasites.
                                                                                                                             (COLD-) COLD SPRING HARBOR LAB
                                                                                                                                                                            WPI; 1997-099928/09
                                  WO9640868-A1
                                                                                                       17-JUN-1995;
                                                         19-DEC-1996.
                                                                                                                                                  Greider C,
           Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADF93862;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 133
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADF93862
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           셤
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present sequence is a PCR primer for the RNA component of a non-human mammal telomerase (RCT), cDNA. The RCT can be used in the recombinant prodn. of an active telomerase mol., capable of adding sequences to chromosomal DNA telomers, and in the synthesis of mutant sequences for the detection of mutant mammalian telomerase RNA component polynucleotides. The RCT may also be used in the identification of telomerase modulating agents, and in the detection of telomerase related, or neoplastic conditions in a patient. Polynucleotides of at least 25 consecutive nucleotides identical, or complementary to the RCT sequence linked to heterologous transcriptional regulatory sequences, can be used for the gene therapy of human diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cancer; eukaryotic parasite; hTR; vertebrate telomerase; yeast; protozoa; tumour; antibody; polymerase chain reaction; ss.
                                                                                                                                                                                                                                                                                                                                                                                          mammalian telomerase, esp. human - useful in identifying
                                                                                                      RNA component; telomerase; recombinant production; synthesis; mutant; detection; mammalian; identification; modulating agent; neoplastic condition; transcriptional regulatory sequence; gene therapy; disease; polymerase chain reaction; PCR primer; non-human; se.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                 RNA component of non-human mammal telomerase cDNA PCR primer R7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               .
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4.7%; Score 21; DB 1; Length 21; 100.0%; Pred. No. 1.2e+02; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 21 BP; 5 A; 5 C; 11 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                              Funk W, Andrews WH;
                                                                                                                                                                                                                                                                                                                                                                                           RNA component of mammalian telomerase, esp. e.g. candidate telomerase-modulating agents
                                                                                                                                                                                                                                                                                                                                                                                                                               Example 9; Page 81; 114pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     184 TGCTGGCCGGTTCGCCCTCC 204
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human telomerase PCR 3'-primer R7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TGCTGGCCCGTTCGCCCCTCC 1
         AAT10301 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                       94US-00330123.
95US-00472802.
95US-00482115.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAT58813 standard; DNA; 21 BP
                                                                                                                                                                                                                                     95WO-US008530
                                                                                                                                                                                                                                                              94US-00272102
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                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                              Villeponteau B, Feng J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21; Conservative
                                                                                                                                                                                                                                                                                                                                                                     WPI; 1996-097581/10.
                                                                                                                                                                                                                                                                                                                       (GERO-) GERON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
                                                                                                                                                                                       WO9601835-A1.
                                                                                                                                                                                                                                    06-JUL-1995;
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                                                         10-SEP-1996
                                                                                                                                                                                                                                                             07-JUL-1994;
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07-JUN-1995;
                                                                                                                                                                                                                                                                                               07-JUN-1995
                                                                                                                                                                                                              25-JAN-1996
                                                                                                                                                               Synthetic.
                                  AAT10301;
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AAT10301/c
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The present sequence represents PCR 3'-primer R7 used for amplifying the human telomerase (hTR). The RNA and DNA can be used in hybridisation cassays to detect or quantify telomerase activity in cells, tissue or fluid samples, e.g. for diagnosis of eukaryotic parasites (yeast and protozoa) or tumours. It is also useful as primers for amplification or assays. The truncated or recombinant vertebrate telomerase is used the charapeutically to increase telomerase activity (also as reagents in the screening assay) while the RNA or other inhibitors such as antisense of therapeutically to reduce such activity. Typical applications are initiation/restoration of activity to cause senescence or to prevent immortalisation of cells in tumours or parasites. The DNA is also used to reduce recombinant telomerase, which can then be used conventionally to craise antibodies for diagnostic detection of telomerase. Detecting telomerase allows early diagnosis of tumour or infection, before clinical signs signs annifest. Telomerase inhibitors directed against e.g. Trypanosoma should cause fewer side effects than drag currently used to treat such infections. The DNA encodes those parts of hTR RNA assential for activity con the endogenous RNA component
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 21 BP; 5 A; 5 C; 11 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          184 TGCTGGCCCGTTCGCCCTCC 204
Example 5; Page 32; 48pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADF93862 standard; RNA; 21 BP
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Best Local Similarity
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Human TERT siRNA, SEQ ID 594.
                                                                                                                                                                                                                                           4.7%;
81.0%;
                         11-FEB-2003; 2003WO-US004088
                                                                          (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                          ADF93867 standard; RNA; 21
                                                                                  Mcswiggen J, Beigelman L;
                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                       Query Match
Best Local Similarity 81.0%
                                                                                           WPI; 2003-689777/65.
        WO2003070742-A1
                                                                                                                                                                                                                                                                                                            26-FEB-2004
                 28-AUG-2003.
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                                                                                                                                                                                                                                                                                  RESULT 134
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1D ADF95

XX

AC ADF95

AC ADF95

DT 26-F1

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Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering nucleic acid; siNA; short interfering RNA; slRNA; double-stranded RNA; micro-RNA; miRNA; short interfering RNA; slRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
                                                                                                                                                                                                                                                                                                                                                                               11-FEB-2003; 2003WO-US004088.
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                                                                                                                                                                                                                                                                                     WO2003070742-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        29-AUG-2002;
05-SEP-2002;
09-SEP-2002;
                                                                                                                                                                                                                                            Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15-JAN-2003;
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                                                                                                                                                                                                                                                                                                                                                                                                                             20-FEB-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      11-MAR-2002;
                                                                                                                                                                                                                                                                                                                                   28-AUG-2003
                                                                                                                                                                                                                   Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short hairpin RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (shRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, can evertor or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA, and vectors that express siNA. The siNAs are expensed expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents a siRNA targeted to the human TERT mRNA transcript.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                invention relates to short interfering nucleic acids (siNA) which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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Pred. No. 1.2e+02;
4; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 21 BP; 2 A; 7 C; 8 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 3; SEQ ID NO 589; 145pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          397 GCGCCGCGATTCCCTGAGCTG 417
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GCGCCCGAUUCCCUGAGCUG 21
                                                                                                                                                                                      20-FEB-2002; 2002US-0358580P.

11-MAR-2002; 2002US-0353124P.

06-JUN-2002; 2002US-0386782P.

17-JUL-2002; 2002US-0396600P.

29-AUG-2002; 2002US-0406784P.

05-SEP-2002; 2002US-0408378P.

09-SEP-2002; 2002US-0409378P.

15-JAN-2003; 2003US-0449129P.
         Homo sapiens.
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2002US-0363124P. 2002US-0386782P. 2002US-039660P. 2002US-0406784P. 2002US-0408378P. 2002US-0409293P.

2003US-0440129P

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The invention relates to short interfering nucleic acids (sinA) which downregulate expression of the one or more telomerase genes by RNA interference. The sinAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and an intense oligonucleotide. Specifically, the sinAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short interfering RNA (siRNA). The sinAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of sinA; conjugates and/or complexes of sinA, and vectors that express sinA. The sinAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restencisis, infectious diseases (specifically protozoal), transplant rejection, or autoimmume or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, ADS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     genetic engineering, pharmacogenomics, studying gene function, and gene
mapping (e.g., of single nucleotide polymorphisms). The present sequence
represents a siRNA targeted to the human TERT mRNA transcript.
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                                                                                                                                                                                        New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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                                                                                                                                                                                                                                                                                                                              Example 3; SEQ ID NO 594; 145pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             395 GCGCGCGCGATTCCCTGAGC 415
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Mcswiggen J, Beigelman L;
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Best Local Similarity 100.
Matches 21; Conservative
                                                                                              WPI; 2003-689777/65.
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Cytostatic, vasotropic, protozoacide, immunosuppressive, dermatological; neuroprotective, anti-HIV; ophthalmological, antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
                      ADF93860 standard; RNA; 21 BP
                                                                                        Human TERT siRNA, SEQ ID 587,
                                                                                                                                                                                                                                                                                                                        2002US-0358580P
2002US-0363124P
                                                                                                                                                                                                                                                                                                                                                        17-JUL-2002; 2002US-0396600P
29-AUG-2002; 2002US-0406784P
05-SEP-2002; 2002US-0408378P.
                                                                                                                                                                                                                                                                                                                                                                                           09-SEP-2002; 2002US-0409293P
15-JAN-2003; 2003US-0440129P
                                                                                                                                                                                                                                                                                                 11-FEB-2003; 2003WO-US004088
                                                                                                                                                                                                                                                                                                                                              2002US-0386782P
                                                                                                                                                                                                                                                                                                                                                                                                                             (RIBO-) RIBOZYME PHARM INC.
                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mcswiggen J, Beigelman L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-689777/65.
                                                                                                                                                                                                                                                    WO2003070742-A1.
                                                                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                                                                         20-FEB-2002;
                                                                                                                                                                                                                                                                                                                                    11-MAR-2002;
06-JUN-2002;
                                                                 26-FEB-2004
                                                                                                                                                                                                                                                                           28-AUG-2003
                                                                                                                                                                                                                  Synthetic
                                            ADF93860;
RESULT 135
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonuclectide and an interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short interfering RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonuclectides, and can be chemically synthesised, can contain deoxyribonuclectides, and can be chemically synthesised, can contain deoxyribonuclectides, and can be chemically synthesised, can contain deoxyribonuclectides, and can be chemically synthesised.

Sexpressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AlbS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug creening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence mapping (e.g., of single nucleotide polymorphisms). The present sequence

Example 3; SEQ ID NO 587; 145pp; English.

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Sequence 21 BP; 2 A; 10 C; 3 G; 0 T; 6 U; 0 Other;
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA committeernee. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and antiense regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairbin RNA (siRNA). The siNAs can be unmodified or chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, can conduct a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be
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                                                                        Gaps
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Score 21; DB 1; Length 21;
Pred. No. 1.2e+02;
6; Mismatches 0; Indels
   Length 21;
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                                                                                                                                                138 CTGCCGCCTTCCACCGTTCAT 158
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2002US-0386782P.
2002US-0396600P.
2002US-0406784P.
2002US-0408378P.
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4.7%;
ilarity 71.4%;
Conservative
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   Query Match
Best Local Similarity
Matches 15; Conserv
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XX AAC ADF9
XX ADF9

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used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, es; multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleoide polymorphisms). The present sequence represents a siRNA targeted to the human TERT mRNA transcript.
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Pred. No. 1.2e+02;
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                                                                                                                                                                                                                                                                                                                                    Sequence 21 BP; 4 A; 13 C; 3 G; 0 T; 1 U; 0 Other;
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Best Local Similarity 100.0%; Pred. No. 1.2s
Matches 21; Conservative 0; Mismatches
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11-MAR-2002; 2002US-0363124P.
06-JUN-2002; 2002US-0386782P.
17-JUJ-2002; 2002US-0396600P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0409378P.
15-JAN-2003; 2003US-0401229P.
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IID ADF93966/6/
AAC ADF93

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an antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (shRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also can contain deoxyribonucleotides, and can be chemically synthesised, can complexe to kite for the in vitro or in vivo delivery of siNA, conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue cyplants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically cortozal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug cortening, disagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents a siRNA targeted to the human TERT mRNA transcript.
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2002US-0408378P.
2002US-0409293P.
2003US-0440129P.
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2002US-0363124P.
2002US-0386782P.
2002US-0396600P.
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06-JUN-2002; 2
17-JUL-2002; 2
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and antisense cregions, or alternatively are assembled from a sense oligonuclectide and an antisense oligonuclectide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short contain deoxyribonuclectides, and can be unmodified or chemically synthesised, can contain deoxyribonuclectides, and can be chemically synthesised. The invention also crelates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express sincells, tissue cexplants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, genetic engineering, pharmacogenomics, studying gene function, and gene capentic engineering, pharmacogenomics, studying gene function, and gene capentic engineering, pharmacogenomics, studying gene function, and gene cepresents a siRNA targeted to the human TERT mRNA transcript.
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  New short interfering nucleic acid downregulates expression of the
                       telomerase gene useful e.g. for treatment and diagnosis of cancer.
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81.0%; Pred. No. 1.2e+02;
tive 4; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 21 BP; 2 A; 3 C; 12 G; 0 T; 4 U; 0 Other;
                                                                    Example 3; SEQ ID NO 586; 145pp; English.
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; 2002US-0363124P.
; 2002US-0386782P.
; 2002US-0396600P.
; 2002US-0406784P.
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nes 17; Conservative
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06-JUN-2002;
17-JUL-2002;
29-AUG-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
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셤
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The invention relates to short interfering nucleic acids (sINA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (shRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue captains (e.g., by ex vivo gene therapy), or in grafts and transplants or the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically sclerosis), lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, disgnosis, therappeutic target identification and validation, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, disgnosis, therappeutic target identification and validation, shin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, disgnosis, therappeutic target identification and validation, appending (e.g., of single multiple polymorphisms). The present sequence mapping (e.g., of single multiple polymorphisms). The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; encorpotective; anti-HIV; ophthalmological; anti-HIV; enclambacold; antiantic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; mitcor.RNA; miRNA; short interfering RNA; siRNA; double-stranded RNA; micro.RNA; miRNA; short hairpin RNA; shoression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              charmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
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                                                                                                                                                                                                                                                    New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4.7%; Score 21; DB 1; Length 21; 95.2%; Pred. No. 1.2e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 21 BP; 6 A; 9 C; 5 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1; Mismatches
                                                                                                                                                                                                                                                                                                                            Example 3; SEQ ID NO 588; 145pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            285 ACCCACTGCCACCGCGAAGAG 305
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 ACCCACUGCCACCGCGAAGAG 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADF93865 standard; RNA; 21 BP
05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409293P.
15-JAN-2003; 2003US-0440129P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human TERT siRNA, SEQ ID 592.
                                                                                                    (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     26-FEB-2004 (first entry)
                                                                                                                                                     Mcswiggen J, Beigelman L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sest Local Similarity 95.2
Atches 20; Conservative
                                                                                                                                                                                                   WPI; 2003-689777/65.
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ADF93865/c
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epilepsy, dementia; huntington's;

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Alzheimer's; Parkinson's; epilepsy; dementia; huntin
amyotrophic lateral sclerosis; gene therapy; ss; hTR
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                                                                         WO2003074654-A2
                                                                                                                                                                20-FEB-2002;
                                                                                                                                                                                           06-JUN-2002;
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05-SEP-2002;
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                                                                                                      12-SEP-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 142
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                                                                                                                                                                                                                                                                                                                                                                   The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense or regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short correlates oligonucleotides, and can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, can complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, itssue captaints or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically proteozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AlDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug correnting, diagnosis, lupus erythematosus, AlDS, macular degeneration, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence correpresents a siRNA targeted to the human TERT mRNA transcript.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   antiarteriosclerotic; neuroprotective; nootropic; antiparkinsonian; anticonvulsant; pulmonary disease; restenosis; atherosclerosis;
                                                                                                                                                                                                                                                                                                 New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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Pred. No. 1.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     double-stranded short interfering nucleic acid; siNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 21 BP; 5 A; 4 C; 11 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                               Example 3; SEQ ID NO 592; 145pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GCCTGCCGCCTTCCACCGTTC 156
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GCTGCCGCCTTCCACCGTTC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Match 4.7%; Sco:
Local Similarity 100.0%; Pr
                                                                         20-FEB-2002; 2002US-0358580P.

11-MAK-2002; 2002US-0363124P.

06-JUN-2002; 2002US-0386782P.

17-JUL-2002; 2002US-0396600P.

29-AUG-2002; 2002US-0406784P.

05-SEP-2002; 2002US-0408378P.

09-SEP-2002; 2002US-0408378P.
                                              11-FEB-2003; 2003WO-US004088
                                                                                                                                                                                15-JAN-2003; 2003US-0440129P
                                                                                                                                                                                                             (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADG30031 standard; RNA; 21
                                                                                                                                                                                                                                           Mcswiggen J, Beigelman L;
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                                                                                                                                                                                                                                                                       WPI; 2003-689777/65.
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                 28-AUG-2003
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The invention relates to a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene comprisiang one or more chemical modifications and each strand of the double-stranded siNA comprises about 21 nucleotides. The siNA of the invention demonstrates antiarteriosclerotic, neuroprotective, noctropic, antiparkinsonian and anticonvulsant activities and may be useful for down-regulating the expression of an endogenous mammalian target gene and therefore in the treatment of any disease or condition that responds to modulation of gene expression or activity in a cell, tissue or organism. The disease or condition may include pulmonary diseases such as restenosis, atherosclerosis, Alzheimer's disease, Parkinson's disease, epillapsy, dementia, huntington's disease or amyotrophic lateral sclerosis. Furthermore, the sinA may be utilised for gene therapy applications. The current sequence is that of the siNA RNA RNA RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New double-stranded short interfering nucleic acid molecule, useful for down-regulating the expression of an endogenous mammalian target gene or for treating diseases that respond to modulation of gene expression or activity.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   .
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Seguence 21 BP; 2 A; 7 C; 8 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 24; SEQ ID NO 597; 593pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mcswiggen J, Beigelman L, Chowri
Jamison S, Usman N, Thompson J;
                                                                                                                                                                                                                                                                                                                                                                                                                         (SIRN-) SIRNA THERAPEUTICS INC
                                                                                                                                                                                                 2002US-0406784P.
2002US-0408378P.
2002US-0409293P.
                                                                                                                      2002US-0363124P.
2002US-0386782P.
20-FEB-2003; 2003WO-US005028
                                                                                                                                                                                                                                                                                                                                     15-JAN-2003; 2003US-0440129P
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nes 17; Conservative
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The invention relates to a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene comprising one or more chemical modifications and each strand of the double-stranded siNA comprises about 21 nucleotides. The siNA of the invention demonstrates antiarteriosalerotic, neuroprotective, nootropic, antiparkinsonian and anticonvulsant activities and may be useful for down-regulating the expression of an endogenous mammalian target gene and therefore in the treatment of any disease or condition that responds to modulation of gene expression or activity in a cell, tissue or organism. The disease or condition may include pulmonary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New double-stranded short interfering nucleic acid molecule, useful for down-regulating the expression of an endogenous mammalian target gene or for treating diseases that respond to modulation of gene expression or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       diseases such as restenosis, atherosclerosis, Aîzheimer's disease, Parkinson's disease, epilepsy, dementia, huntington's disease or amyotrophic lateral sclerosis. Purthermore, the siNA may be utilised for gene therapy applications. The current sequence is that of the siNA RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      antiarteriosclerotic; neuroprotective; nootropic; antiparkinsonian; anticonvulsant; pulmonary disease; restenosis; atherosclerosis;
 Alzheimer's; Parkinson's; epilepsy; dementia; huntington's; amyotrophic lateral sclerosis; gene therapy; se; hTR.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       double-stranded short interfering nucleic acid; siNA;
                                                                                                                                                                                                                                                                                                                                                                                    Chowrira B, Pavco P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 21 BP; 6 A; 9 C; 5 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 24; SEQ ID NO 596; 593pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                       Thompson J;
                                                                                                                                                                                                                                                                                                                                                (SIRN-) SIRNA THERAPEUTICS INC
                                                                                                                                                                                                                 2002US-0363124P.
2002US-0386782P.
2002US-0406784P.
2002US-0408378P.
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                                                                                                                                                                20-FEB-2003; 2003WO-US005028.
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15-JAN-2003; 2003US-0440129P.
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ID ADG30036 standard; RNA; 21
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                                                                                                                                                                                                                                                                                                                                                                                    Beigelman
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R. Usman N,
                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2003-731676/69.
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Best Local Similarity
Matches 20; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  of the invention
                                                                                        WO2003074654-A2
                                                      Unidentified
                                                                                                                                                                                                                                        06-JUN-2002;
                                                                                                                                                                                                                                                     29-AUG-2002;
05-SEP-2002;
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                                                                                                                            12-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                                                     Jamison S,
                                                                                                                                                                                                                                                                                                                                                                                    Mcswiggen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 activity.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene comprising one or more chemical modifications and each strand of the double-stranded siNA comprises about 21 nucleotides. The siNA of the invention demonstrates antiarteriosclerotic, neuroprotective, nootropic, antiparkinsonian and anticonvulsant activities and may be useful for down-regulating the expression of an endogenous mammalian tharget gene and therefore in the treatment of any disease or condition that responds to modulation of gene expression or activity in a cell, tissue or organism. The disease or condition may include pulmonary diseases such as restenosis, atherosclerosis, Alzheimer's disease, parkinson's disease, epilepsy, dementia, hutington's disease or amyotrophic lateral sclerosis. Furthermore, the siNA may be utilised for gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New double-stranded short interfering nucleic acid molecule, useful for down-regulating the expression of an endogenous mammalian target gene or for treating diseases that respond to modulation of gene expression or
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   dementia; huntington's;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 4.7%; Score 21; DB 1; Length 21; Best Local Similarity 81.0%; Pred. No. 1.2e+02; Matches 17; Conservative 4; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       double-stranded short interfering nucleic acid; siNA;
Alzheimer's; Parkinson's; epilepsy; dementia; huntin
amyotrophic lateral sclerosis; gene therapy; ss; hTR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 21 BP; 2 A; 3 C; 12 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 24; SEQ ID NO 594; 593pp; English.
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man N, Thompson J;
                                                                                                                                                                                                                 2002US-0363124P.
2002US-0386782P.
2002US-0406784P.
2002US-0408378P.
                                                                                                                                                                                                                                                                                                                                              (SIRN-) SIRNA THERAPEUTICS INC
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                                                                                                                                                                                                                                                     29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408376P.
09-SEP-2002; 2002US-0409293P.
                                                                                                                                                                                                                                                                                                              15-JAN-2003; 2003US-0440129P
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                                                                                                                                                                                                                                                                                                                                                                 Mcswiggen J, Belyerman N,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  of the invention
                                                                                      WO2003074654-A2
                                                      Unidentified.
                                                                                                                                                                                                                                      06-JUN-2002;
                                                                                                                                                                                                                     11-MAR-2002;
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Foenaugh K;

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Alzheimer's; Parkinson's; epilepsy; dementia; huntington's; amyotrophic lateral sclerosis; gene therapy; ss; hTR.
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                                                                                                                                                                                                                                                                                                         WPI; 2003-731676/69.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         of the invention
                                                                WO2003074654-A2
                                                                                                                                                                              29-AUG-2002; 2
05-SEP-2002; 2
09-SEP-2002; 2
15-JAN-2003; 2
                                      Unidentified.
                                                                                                                                           20-FEB-2002;
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antiarteriosclerotic; neuroprotective; nootropic; antiparkinsonian;
anticonvulsant; pulmonary disease; restenosis; atherosclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ;
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                                                                                                                                                                                                                                                                        Pavco P, Fosnaugh K;
 Alzheimer's, Parkinson's, epilepsy, dementia, huntington's, amyotrophic lateral sclerosis, gene therapy; ss; hTR.
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Pred. No. 1.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 21 BP; 3 A; 8 C; 8 G; 0 T; 2 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                   Example 24; SEQ ID NO 602; 593pp; English.
                                                                                                                                                                                                                                                                      Beigelman L, Chowrira B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  hTR-targeted sina RNA - SEQ ID 601
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                                                                                                                                                                                                                                                                                     Thompson J;
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11-MAR-2002; 2002US-0363124P.
06-JUN-2002; 2002US-0366782P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409293P.
                                                                                                                                                                                                                                               (SIRN-) SIRNA THERAPEUTICS INC
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                                                                                                                   20-FEB-2003; 2003WO-US005028
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                                                                                                                                                                                                                                                                                    Usman N,
                                                                                                                                                                                                                                                                                                          WPI; 2003-731676/69.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             the invention
                                                                 WO2003074654-A2
                                         Unidentified
                                                                                                                                                                                                                                                                         Mcswiggen J,
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                                                                                           12-SEP-2003
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                                                                                                                                                                                                                                                                                     Jamison S,
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The invention relates to a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene comprishing one or more chemical modifications and each strand of the double-stranded siNA comprises about 21 nucleotides. The siNA of the invention demonstrates antiarteriosclerotic, neuroprotective, noctropic, antiparkinsonian and anticonvulsant activities and may be useful for down-regulating the expression of an endogenous mammalian target gene and therefore in the treatment of any disease or condition that responds to modulation of gene expression or activity in a cell, tissue or organism. The disease or condition may include pulmonary diseases such as restenosis, atherosclerosis, Alzheimer's disease, Parkinson's disease, epilapsy, dementia, huntington's disease, epilabsy, dementia, huntington's disease or amyotrophic lateral sclerosis. Furthermore, the sinA may be utilised for gene therapy applications. The current sequence is that of the sinA RNA RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New double-stranded short interfering nucleic acid molecule, useful for down-regulating the expression of an endogenous mammalian target gene or for treating diseases that respond to modulation of gene expression or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 24; SEQ ID NO 601; 593pp; English.
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Jamison S, Usman N, Thompson J;
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                                                                                                                                                                                                                                                                                                                               (SIRN-) SIRNA THERAPEUTICS INC
2002US-0363124P.
2002US-0386782P.
                                                                                     2002US-0406784P.
2002US-0408378P.
2002US-0409293P.
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or

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Alzheimer's; Parkinson's; epilepsy; dementia; huntington's; amyotrophic lateral sclerosis; gene therapy; ss; hTR.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  of the invention.
                                                           WO2003074654-A2
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Local S...
21;
                                    Unidentified
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 Alzheimer's; Parkinson's; epilepsy; dementia; huntington's; amyotrophic lateral sclerosis; gene therapy; 88; hTR.
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Pred. No. 1.2e+02;
6; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                   Example 24; SEQ ID NO 595; 593pp; English.
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| CUGCCGCCUUCCACCGUUCAU 21
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sman N, Thompson J;
                                                                                                                                 2002US-0358580P.
2002US-0363124P.
2002US-0386782P.
2002US-0406784P.
2002US-040878P.
                                                                                                                                                                                                                                   (SIRN-) SIRNA THERAPEUTICS INC
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                                                                                                            20-FEB-2003; 2003WO-US005028
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Conservative
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                                                                                                                                                                                                                                                Mcswiggen J, Belyerman S. Usman N,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
ses 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   the invention
                                                             WO2003074654-A2
                                      Unidentified
                                                                                                                                                                        29-AUG-2002;
05-SEP-2002;
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                                                                                    12-SEP-2003
                                                                                                                                                                                                                                                                                                                                                          activity.
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The invention relates to a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene comprising one or more chemical modifications and each strand of the double-stranded siNA comprises about 21 nucleotides. The siNA of the invention demonstrates antiarteriosclerotic, neuroprotective, nootropic, antiparkinsonian and anticonvulsant activities and may be useful for down-regulating the expression of an endogenous mammalian target gene and therefore in the treatment of any disease or condition that responds to modulation of gene expression or activity in a cell, tissue or organism. The disease or condition may include pulmonary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New double-stranded short interfering nucleic acid molecule, useful for down-regulating the expression of an endogenous mammalian target gene or for treating diseases that respond to modulation of gene expression or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          diseases such as restenosis, atherosclerosis, Alzheimer's disease, Parkinson's disease, epilepsy, dementia, huntington's disease or amyotrophic lateral sclerosis. Furthermore, the sINA may be utilised for gene therapy applications. The current sequence is that of the sINA RNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 double-stranded short interfering nucleic acid; siNA; antiarteriosclerotic; neuroprotective; nootropic; antiparkinsonian; anticonvulsant; pulmonary disease; restenosis; atherosclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ..
                                                                                                                                                                                                                                                                                                                                                                                            ĸ
                                                                                                                                                                                                                                                                                                                                                                                            Fosnaugh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 21; DB 1; Length 21;
Pred. No. 1.2e+02;
0; Mismatches 0; IndelB
                                                                                                                                                                                                                                                                                                                                                                                            B, Pavco P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 21 BP; 4 A; 13 C; 3 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 24; SEQ ID NO 599; 593pp; English.
                                                                                                                                                                                                                                                                                                                                                                                            Chowrira
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        hTR-targeted sina RNA - SEQ ID 600.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             22
                                                                                                                                                                                                                                                                                                                                                                                                                             Thompson J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4.' 100.0%; Pre
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          н
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                                                                                                                                                                                                                                                                                                                               (SIRN-) SIRNA THERAPEUTICS INC
                                                            2002US-0358580P.
2002US-0363124P.
2002US-0386782P.
2002US-0406784P.
2002US-0408378P.
                                                                                                                                                                                                                          09-SEP-2002; 2002US-0409293P.
20-FEB-2003; 2003WO-US005028
                                                                                                                                                                                                                                                                                                                                                                                            Beigelman L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              51
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADG30034 standard; RNA;
                                                                                                                                                                                                                                                                                                                                                                                  1 J, Bels
G Usman N,
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cardiovascular disease; immune disease; rheumatoid arthritis;
              angiogenic abnormality
                                                                                                                                                                                                    WPI; 2004-553738/53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO2004063380-A1
                                                           WO2004063380-A1
                                                                                                                                                                                                                                                               abnormalities.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29-JUL-2004
                                                                                 29-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
                                     Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADQ94243;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                             Series
                                                                                                                                                                               Ψu Χ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Loc
Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene comprising one or more chemical modifications and each strand of the double-stranded siNA comprises about 21 nucleotides. The siNA of
                                                                                                                                                                                                                                                                                                                                                                                                                                        the invention demonstrates antiarteriosclerotic, neuroprotective, nootropic, antiparkinsonian and anticonvulsant activities and may be useful for down-regulating the expression of an endogenous mammalian target gene and therefore in the treatment of any disease or condition that responds to modulation of gene expression or activity in a cell, tissue or organism. The disease or condition may include pulmonary laiseases such as restenoosis, atherosclerosis, Albrahimer's disease, Parkinson's disease, epilepsy, dementia, huntington's disease or amyotrophic lateral sclerosis. Furthermore, the siNA may be utilised for gene therapy applications. The current sequence is that of the siNA RNA
                                                                                                                                                                                                                                                                                                            New double-stranded short interfering nucleic acid molecule, useful for down-regulating the expression of an endogenous mammalian target gene or for treating diseases that respond to modulation of gene expression or activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ss; antiarthritic; antirheumatic; cardiovascular; cytostatic; gene therapy; adeno-associated virus; RNAi; RNA interference; short interfering RNA; gene silencing; short hairpin RNA; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ..
0
                                                                                                                                                                                                                                                     Fosnaugh K;
   dementia; huntington's;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 21; DB 1; Length 21;
Pred. No. 1.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Indels
  Alzheimer's; Parkinson's; epilepsy; dementia; huntin
amyotrophic lateral sclerosis; gene thexapy; ss; hTR
                                                                                                                                                                                                                                                     Pavco P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 21 BP; 5 A; 4 C; 11 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                        Example 24; SEQ ID NO 600; 593pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Short hairpin RNA oligonucleotide SHRNA12.
                                                                                                                                                                                                                                                     Beigelman L, Chowrira B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GCCTGCCGCCTTCCACCGTTC 156
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4.7%; Scor.
100.0%; Pred
0; N
                                                                                                                                                                                                                                                                  Thompson J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GCCTGCCGCCTTCCACCGTTC 1
                                                                                                                                                                                                                              (SIRN-) SIRNA THERAPEUTICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ВР
                                                                                                                                             2002US-0363124P.
2002US-0386782P.
2002US-0406784P.
                                                                                                                                                                                2002US-0408378P.
2002US-0409293P.
                                                                                                            20-FEB-2003; 2003WO-US005028
                                                                                                                                  2002US-0358580P
                                                                                                                                                                                                        15-JAN-2003; 2003US-0440129P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADQ94244 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                  Usman N,
                                                                                                                                                                                                                                                                                        WPI; 2003-731676/69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                              WO2003074654-A2
                                                                                                                                                        06-JUN-2002;
29-AUG-2002;
05-SEP-2002;
09-SEP-2002;
                                                                                                                                                                                                                                                     Mcswiggen J,
Jamison S, I
                                       Unidentified
                                                                                                                                   20-FEB-2002;
                                                                                                                                              11-MAR-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21-OCT-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21;
                                                                                     12-SEP-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 149
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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    셤
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The invention relates to an adeno-associated virus (AAV) that carries a specific RNAi (RNA interference) nucleotide fragment comprising components of the outer shell of a recombinant AAV, a specific RNAi nucleotide fragment, and a control element for regulating the transcription and expression of the RNAi nucleotide fragment carried by encapsulation within the outer shell of a recombinant AAV. The recombinant viruses are useful for inducing RNAi pathway and gene therapy in treating e.g. cancer, cardiovascular diseases, immune diseases like rheumatoid arthritis and angiogenic abnormalities. This sequence corresponds to a short hairpin RNA (shRNA) oligonucleotide used in the
                                                                                                                                                                                                                                                                                                                                                                                            recombinant adeno-associated viruses useful for inducing RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                  interference pathway and for gene therapy in treating e.g. cancer, cardiovascular diseases, rheumatoid arthritis and angiogenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        se; antiarthritic; antirheumatic; cardiovascular; cytostatic; gene therapy; adeno-associated virus; RNAi; RNA interference; short interfering RNA; gene silencing; short hairpin RNA; cancer; cardiovascular disease; immune disease; rheumatoid arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ó,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ch 4.7%; Score 21; DB 1; Length 21; 1 Similarity 100.0%; Pred. No. 1.2e+02; 21; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 21 BP; 4 A; 4 C; 8 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Short hairpin RNA oligonucleotide SHRNA11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 8; SEQ ID NO 20; 53pp; Chinese.
                                                                                                                                                                                                                                      Hou Y;
                                                                                                                                                      (AGTC-) AGTC GENE TECHNOLOGY CO LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           300 GAAGAGTTGGGCTCTGTCAGC 320
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GAAGAGTTGGGCTCTGTCAGC 21
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                                                                                                                                                                                                                                          Ľu X,
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07-NOV-2003; 2003WO-CN000939.
                                                                           07-NOV-2002; 2002CN-00149319
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               angiogenic abnormality
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Disclosure; Page 24; 79pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            27-OCT-1994;
13-FEB-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                07-JUL-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-JAN-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAT11035;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 152
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAT11035,
원
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   8
                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                          The invention relates to an adeno-associated virus (AAV) that carries a specific RNAi (RNA interference) nucleotide fragment comprising components of the outer shell of a recombinant AAV, a specific RNAi nucleotide fragment, and a control element for regulating the transcription and expression of the RNAi nucleotide fragment carried by encapsulation within the outer shell of a recombinant AAV. The recombinant viruses are useful for inducing RNAi pathway and gene therapy in treating e.g. cancer, cardiovascular diseases, immune diseases like rheumatoid arthritis and angiogenic abnormalities. This sequence corresponds to a short hairpin RNA (shRNA) oligonucleotide used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          rRNA processing; RNA methylation site selection; pseudouridine formation; HACA-enoRNA; telomerase RNA; cancer; tumour; cell proliferation; trypanosome infection; RNNi-mediated degradation; neo gene; siRNA; small interfering RNA; TER; human; telomerase; RNAi; selection marker.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               88; snoRNA; small nuclear RNA; box H/ACA; mRNA splicing; mRNA processing;
                                                                                          Series of recombinant adeno-associated viruses useful for inducing RNA interference pathway and for gene therapy in treating e.g. cancer, cardiovascular diseases, rheumatoid arthritis and angiogenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New isolated small nuclear RNA (snoRNA) polynucleotides, useful for inducing RNAi-mediated degradation of snoRNA or for treating diseases associated with activity of small nuclear RNA, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                                                                             4.7%; Score 21; DB 1; Length 21; 100.0%; Pred. No. 1.2e+02; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                         Sequence 21 BP; 3 A; 8 C; 3 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       siRNA sequence used for RNA inhibition.
                                                                                                                                                      Example 8; SEQ ID NO 19; 53pp; Chinese.
                       (AGTC-) AGTC GENE TECHNOLOGY CO LTD.
                                                                                                                                                                                                                                                                                                                                                                                               143 GCCTTCCACCGTTCATTCTAG 163
                                                                                                                                                                                                                                                                                                                                                                                                            1 GCCTTCCACCGTTCATTCTAG 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADT86999 standard; RNA; 21 BP
07-NOV-2002; 2002CN-00149319.
                                              Ľu x,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         04-FEB-2004; 2004WO-IL000108
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                         21; Conservative
                                              Ma X,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (UYBA-) UNIV BAR-ILAN
                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
                                                                     WPI; 2004-553738/53.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO2004069148-A2.
                                               Wu X, Dong X,
                                                                                                                                abnormalities.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16-DEC-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Michaeli S;
                                                                                                                                                                                                                                                                                                    invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADT86999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 151
                                                                                                                                                                                                                                                                                                                                                                          Matches
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The present invention provides the method for downregulating snokkh.

The present invention provides the method for downregulating snokkh.

The method can be used to treat the diseases associated with the activity of small nuclear RNA Small nuclear RNA molecules are the important regulators of gene expression. They participate in mRNA splicing, mRNA and rRNA processing, RNA methylation site selection and pseudouridine for formation (box HAACA-snokNA). The telomerase RNA is an important nuclear RNA which serves as a template for telomerase replication. It contains box HAACA like domain which confers the functional localisation of this CNA which serves as a template for telomerase replication. It contains box HAACA like domain which confers the functional localisation of this economic contain active telomerase which contributes to cell proliferation. It has been found that the trypanosome infection is associated with the parasite's snoRNA expression. The polynucleotides of the invention are useful for inducing RNA-mediated degradation of a small nuclear RNA. They are useful for down regulating snoRNA molecules or box H/ACA-containing RNA mediated degradation of the cancer cells can be inhibited by a polynucleotide sequence which includes two additional coding sequences of novel RNA termed as h2 and h3. It has also observed that the snoRNA-2 guided methylation on the 5.8FNA. The results in decreased snoRNA-2 guided methylation on the S.8FNA. The results in decreased snoRNA-2 guided methylation on the S.8FNA. The production of siRNA (small interfering RNA) which can produce in both nucleus and cytoplasm. The invention suggests that the silencing of snoRNA-2 silencing of snoRNA-2 silencing of snoRNA-2 transcript RNA which can produce in both remarker some and cytoplasm. The invention suggests that the silencing of snoRNA-2 silencing snot unique to snoRNA cluster-2. It is also contained to a selection marker gene in reverse orientation. It also transcription of a snoRNA-2 shear snoRNA-2 shear snoRNA-2 shear snoRNA-2 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             transcription of a specific strand of the nucleotide. The presented nucleotide sequence is the siRNA sequence which was used for snRNA
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The present invention provides the method
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94US-00330123.
95US-00387524.
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US09436060A-16.rng.sl

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Villeponteau B,
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                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                              AAT10286;
                                                                                                                                                                                                                                                                                         RESULT 154
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                                                                                                                                             The RNA components of (non) human mammalian telomerase(s) especially from mouse, rat and chinese hamster are all claimed. Antisense oligonucleotides can be used to block the activity of the telomerase; probes and primers can be used in detection; vectors and host cells transformed with the isolated telomerase genes can be used for production of telomerases; RNA and DNA ribozymes and triplex forming oligonucleotides directed against the telomerase genes can be used to therapeutically as can plasmids. A mouse which lacks the telomerase gene therapeutically as can plasmids. A mouse which lacks the telomerase gene the role it plays in immortalisation. The antisense oligonucleotide is synthesised as a 2-0-methyl RNA oligonucleotide and is more resistant to hydrolysis than unmodified RNA oligonucleotides (See AATI1032-35)
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                                                                                                RNA components of (non)human mammalian telomerase(s) - useful in studying
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     mammal; antisense; triplex forming oligonucleotide; plasmid;
                                                  Marhuenda MA;
                                                                                                                                                                                                                                                                                                                     Gaps
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                                                                                                                                                                                                                                                                                               4.4%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 1.3e+02; ive 0; Mismatches 0; Indels
                                                  Greider C,
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                                                                                                                                                                                                                                                                           Sequence 20 BP; 4 A; 12 C; 3 G; 1 T; 0 U; 0 Other;
                                                 Funk W,
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                                                                                                           cell senescence and immortalisation.
                                                                                                                            Disclosure; Page 23; 85pp; English
                                                 Feng J,
                                                                                                                                                                                                                                                                                                                                      2 GGTTGCGGAGGGTGGGCCTG 21
                                                                                                                                                                                                                                                                                                                                               20 GGTTGCGGAGGGTGGGCCTG 1
                    SPRING HARBOR LAB
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94US-00330123.
95US-00387524.
95US-00485778.
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 95US-00485778,
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probe; primer; ribozyme; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                          Villeponteau B;
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                                                                                                                                                                                                                                                                                                                   20; Conservative
                                                                             WPI; 1996-097428/10
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                            (GERO-) GERON CORP
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07-JUN-1995;
                   (COLD-) COLD
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13-FEB-1995;
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                                                                                                                                                                                                                                                                                               Query Match
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mouse, rat and chinese hamster are all claimed. Antisense of objective to be objected to block the activity of the telomerase; probes and primers can be used to block the activity of the telomerase; probes and primers can be used to lact cells transformed with the isolated telomerase genes can be used for production of telomerases; RNA and DNA ribozymes and triplex forming oligonucleotides directed against the telomerase genes can be used therapeutically as can plasmids. A mouse which lacks the telomerase gene therapeutically as can plasmids. A mouse which lacks the telomerase gene the role it plays in immortalisation. The antisense oligonucleotide is synthesised as a 2-0-methyl RNA oligonucleotide and is more resistant to hydrolysis than unmodified RNA oligonucleotides (See AATI1032-35)
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RNA components of (non)human mammalian telomerase(s) - useful in studying cell senescence and immortalisation.
                                                                                                                                                                          The RNA components of (non) human mammalian telomerase(s) especially from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RNA component of mammalian telomerase, esp. human - useful in identifying e.g. candidate telomerase-modulating agents.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RNA component; mammalian; telomerase; antisense oligonucleotide; triple helix; inhibition; neoplastic; cells; activity; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1; Length 20;
1.3e+02;
thes 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 7 A; 3 C; 5 G; 0 T; 5 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4.4%; Score 20; DB 100.0%; Pred. No. 1.3
                                                                                                         Disclosure; Page 23; 85pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure, Page 38; 114pp; English.
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94US-00330123.
95US-00472802.
95US-00482115.
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ID AAT10286 standard; DNA; 20
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Best Local Similarity 100.0
Matches 20; Conservative
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vaccine; cancer; antisense oligonucleotide; ss.
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              셤
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present sequence is a RNA component of mammalian telomerase, antisense oligonucleotide, which can be used, along with triple helix forming sequences, to inhibit telomerase activity in cells, esp.
                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Antisense oligonucleotide 14ab for human telomerase RNA component
                                                                                                                                                                                                                                                                                                                                                                                                      RNA component; mammalian; telomerase; antisense oligonucleotide;
triple helix; inhibition; neoplastic; cells; activity; ss.
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                                                                                                                                                                                                                                                                                                                                                                RNA component of mammalian telomerase antisense oligo Tel-AU
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            Length 20;
                                                     0; Indels
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            Score 20; DB 1; Le
Pred. No. 1.3e+02;
0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  e.g. candidate telomerase-modulating agents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 38; 114pp; English.
4.4%; Scc...
100.0%; Pred
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                                                                                                                     20 TTTGTCTAACCCTAACTGAG 1
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                                                                                             41 TITGICIAACCCIAACTGAG
                                                                                                                                                                                                                                        BP
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94US-00330123.
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95US-00482115.
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AAT10289 standard; DNA; 20
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Best Local Similarity 100.
Matches 20; Conservative
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                                                       Conservative
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                                  Similarity
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07-JUN-1995;
07-JUN-1995;
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          Query Match
Best Local (
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                                    Best Loc
Matches
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XXX

AAC AAT11

XXX AAT11

XXX AAT11

XXX COS Syntl

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ID AAV7
XX AC AAV7
XX DT 15-F
XX Anti
XX Huma
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The present sequence represents an antisense oligonucleotide directed against the human telomerase RNA component gene sequences. The oligonucleotide can be used as an affinity agent in the methods of the invention, which are used to purify human telomerase. The methods of the the use of several sequential steps, including the use of two matrixes that bind molecules bearing negative charges, a matrix that binds molecules bearing negative charges, an affinity purification step and a size separation. Telomerase is a particular target of anticancer therapies, and is useful in assays for characterizing (pre) cancerous cells. Telomerase can also be used to screen for specific modulators, for biochemical analysis of its activity, and in preparation of antibodies. Fragments of telomerase, or nucleic acid encoding them, are used in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Fragments of telomerase, or nucleic acid encoding them, are used in vaccines, and for treating over expression of telomerase, particularly in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RNA component; human telomerase; antisense oligonucleotide; infection; neuroblastoma; bladder cancer; colon cancer; prostate cancer; contraception; sterilisation; immunosuppression; therapeutic; hTR; immune system down-regulation; anti-inflammatory therapy; ss.
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                                                                                                                                                                                                                                                                     /*tag= a
/note= "biotinylated"
                                                                                                                                                         Location/Qualifiers
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ID AAV41173 Standard; DNA; 20 BP XX
AC AAV41173;
XX
DT 08-OCT-1998 (first entry)
XX
DE RNA component of human telomera
XX
W RNA component; human telomera
XX
KW RNA component; human telomera
XX
Contraception; sterilisation;
XX
Contraception; Sterilisation;
XX
Contraception; Sterilisation;
XX
Contraception; Sterilisation
XX
Contraception; Sterilisation;
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Contraception; Sterilisation; Sterilis
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Best Local Similarity 100.
Matches 20; Conservative
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modified_base
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                                                   Homo sapiens
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Synthetic
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WO9828442-A1
WO9828442-A1
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oligonucleotides to the RNA component of human telomerase (hTR). These
antisense oligonucleotides specifically bybridise to a nucleotide
sequence within an accessible region of the hTR, but that does not
hybridise to sequence within the template region of hTR. These
component of human telomerase in a sample. This is useful for disponate of human telomerase in a sample. This is useful for disponant of human telomerase in a lample. This is useful for disponant of human telomerase in a lample. This is useful for disponant of human telomerase in a cancer patient. The inhibitor cancer),
and providing prognosis for a cancer patient. The inhibitor and providing with transcription of the RNA component, decreasing the half-
interfering with transcription of the RNA component, decreasing the half-
interfering with transcription of the RNA component, decreasing the half-
interfering with transcription of the RNA component into the telomerase holoenzyme, or inhibiting assembly of
the RNA component into the telomerase. These antisense oligonucleotides can
be used for inhibiting telomerase activity in both cultured cells and in
cells in vivo. They can be used in therapeutics for treating or
immunosuppression, and for selectively down-regulating specific branches
inflammatory theraples or for treating infections by, e.g. yeast,
parasites or fungi
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                                                                                                                                                                                                         New polynucleotide(s) anti:sense to human telomerase - used for detecting or inhibiting human telomerase, e.g. for treating cancers, contraception, immuno-suppression or treating infection.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RNA component of human telomerase (hTR) antisense oligo 14ab.
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100.0%; Pred. No. 1.3e+02;
ive 0; Mismatches 0; Indels
                                                                                                                                                         Weinrich SL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 5 A; 8 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                          Kealey JT, Pruzan R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            300 GAAGAGTTGGGCTCTGTCAG 319
                                                                                                                                                                                                                                                               Claim 11; Page 65; 80pp; English
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                                                                 97WO-US023619
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96US-00770565
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                                                                                                                                                                                   WPI; 1998-377670/32.
                                                                                                                                  (GERO-) GERON CORP.
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                                                                19-DEC-1997;
                                                                                          20-DEC-1996;
20-DEC-1996;
                WO9828442-A1
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                                                                                                                                                         Kim NW,
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Sequences shown in AAV41169 to AAV41181 represent antisense
oligonucleotides to the RNA component of human telomerase (hTR). These
antisense oligonucleotides specifically hybridise to a nucleotide
sequence within an accessible region of the hTR, but that does not
hybridise to a sequence within the template region of hTR. These
component of human telomerase in a sample. This is useful for diagnosing
cancer (especially neuroblastoma, bladder, colon and prostate cancer),
and providing prognosis for a cancer patient. The inhibitory
cancer (especially neuroblastoma, bladder, colon and prostate cancer),
and providing prognosis for a cancer patient. The inhibitory
coligonucleotides can inhibit the telomerase activity level in a call by
interfering with transcription of the RNA component, decreasing the half-
colisting assembly of telomerase holoenzyme, or inhibiting the epoperase activity of telomerase. These antisense oligonucleotides can
collymerase activity of telomerase. These antisense oligonucleotides can
be used for inhibiting telomerase. These antisense oligonucleotides can
cells in vivo. They can be used in therapeutics for treating or
cells in vivo. They can be used in therapeutics for treating or
cells in vivo. and for selectively down-requiating specific branches
confimunosuppression, and for selectively down-requiating specific branches
confirmancy theraples or for treating infections by, e.g. yeast,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               - used for detecting
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100.0%; Pred. No. 1.3e+02;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                      Pruzan R, Weinrich SL;
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                                                                              97WO-US023619.
                                                                                                                                                             96US-00770564
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Best Local Similarity 100.0
Matches 20; Conservative
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parasites or fungi
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                                                                              19-DEC-1997;
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02-JUL-1998
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04-APR-1997;
02-JUL-1998.
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                                                                                                            Kim NW,
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                                                                                                                                                                                                                                              Sequences shown in AAV41169 to AAV41181 represent antisense
oligonucleotides to the RNA component of human telomerase (hTR). These
antisense oligonucleotides specifically hybridise to a nucleotide
equence within an accessible region of the hTR, but that does not
hybridise to a sequence within the template region of hTR. These
oligonucleotides may specifically be used for detection of an RNA
component of human telomerase in a sample. This is useful for diagnosing
cancer (especially neuroblastom, bladder, colon and prostate cancer),
and providing prognosis for a cancer patient. The inhibitory
oligonucleotides can inhibit the telomerase activity level in a cell by
interfering with transcription of the RNA component, decreasing the half-
life of the telomerase RNA component transcript, inhibiting assembly of
the RNA component into the telomerase holoenzyme, or inhibiting the
polymerase activity of telomerase activity in both cultured cells and in
cells in vivo. They can be used in therapeutics for treating or
preventing cancer, for contraception or sterilisation, for
immunosuppression, and for selectively down-regulating specific branches
of the immune system, e.g. a specific subset of T-cells, in anti-
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                                                                                                                                                                       New polynucleotide(s) anti:sense to human telomerase - used for detecting or inhibiting human telomerase, e.g. for treating cancers, contraception, immuno-suppression or treating infection.
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4.4%; Score 20; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 1.3e+02;
Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                         Wu F, Kealey JT, Pruzan R, Weinrich SL;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            290 CTGCCACCGCGAAGAGTTGG 309
                                                                                                                                                                                                                            Claim 11; Page 65; 80pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20 CTGCCACCGCGAAGAGTTGG 1
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                                                             96US-00770564
                                                                         96US-00770565
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                                                                                                 (GERO-) GERON CORP.
                                    19-DEC-1997;
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20-DEC-1996;
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             02-JUL-1998
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Sequences shown in AAV41169 to AAV41181 represent antisense oligonucleotides to the RNA component of human telomerase (hTR). These antisense oligonucleotides specifically hybridise to a nucleotide sequence within an accessible region of the hTR, but that does not hybridise to a sequence within the template region of hTR. These oligonucleotides may specifically be used for detection of an RNA component of human telomerase in a sample. This is useful for diagnosing cancer (especially neuroblastoma, bladder, colon and prostate cancer), and providing prognosis for a cancer patient. The inhibitory oligonucleotides can inhibit the telomerase activity level in a cell by interfering with transcription of the RNA component, decreasing the half-life of the telomerase RNA component transcript, inhibiting assembly of the RNA component into the telomerase holoenzyme, or inhibiting the polymerase activity of telomerase. These antisense oligonucleotides can be used for inhibiting telomerase activity of telomerase. These antisense oligonucleotides and in
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    used for detecting

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 4.4%; Score 20; DB 1; Length 20; Similarity 100.0%; Pred. No. 1.3e+02; 0; Conservative 0; Mismatches 0; Indel8
                                                                                                                                                                                                                                                              Weinrich SL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 3 A; 3 C; 3 G; 11 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human clone 28-1 telomerase oligonucleotide 14ab.
                                                                                                                                                                                                                                                              Wu F, Kealey JT, Pruzan R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               159 TCTAGAGCAAACAAAATG 178
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 11; Page 65; 80pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20 TCTAGAGCAAACAAAAATG 1
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97WO-US023619
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96US-00770565
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                                                                                                                                                                                                                                                                                                                                         WPI; 1998-377670/32.
                                                                                                                                                                                    (GERO-) GERON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mar
Local Sir.
20; f
                                                                          20-DEC-1996;
20-DEC-1996;
19-DEC-1997;
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associated with overexpression of telomerase e.g. cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 163
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                                                                                                                                                                                                                                                                                                                                                                                                                            Db
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ;
0
                                                                                                                                                                                                                                                                                       This invention describes a novel composition comprising human telomerase having at least 2000-fold (preferably at least 60000-fold) increased relative purity compared with crude extract of cells from adenovirustransformed kidney cell line. The composition is useful for eliciting an immune response in animals and may therefore be used as a vaccine for treating diseases associated with the overexpression of telomerase e.g. cancer. AAZ2105-C22031 represent oligonucleotides used in the isolation of human clone 28-1 which contains a fragment of the human telomerase described in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Telomerase; human; immune response; cancer; vaccine; treatment; disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Compositions comprising human telomerase, useful for treating diseases
                                                                                                                                                                                                 treating diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                  Kealey JT;
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 20; DB 1; Length 20;
Pred. No. 1.3e+02;
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                                                                                                  Pruzan RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Indels
                                                                                                                                                                                                 Compositions comprising human telomerase, useful for treassociated with overexpression of telomerase e.g. cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human clone 28-1 telomerase oligonucleotide oligo 14ab
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 0 A; 9 C; 4 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         a
"5'-biotinylated cytidine"
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                                                                                                  Lichtsteiner SP, Weinrich SL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4.4%; Scor.
100.0%; Pred. No. ...
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                                                                                                                                                                                                                                                           Disclosure, Col 45-46; 34pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         361 AGGCCGCAGGAAGAGGAACG 380
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP
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                          95US-00510736
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 100.
Matches 20; Conservative
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/note=
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                                                                                                                                                            WPI; 1999-590379/50
                                                                (GERO-) GERON CORP.
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                          04-AUG-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
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Vasserot AP;
                                                                                                    Atkinson EM,
Vasserot AP;
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                                                                                                                          Vasserot
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The invention relates to promoter regions from mouse and human telomerase RNA (TR) component genes. The TR gene promoter can be linked to a heterologous gene, especially a gene encoding a cytotoxin, for therapy of cancer, especially neoplasias. The telomerase is necessary for the unrestricted proliferative capacity of many human cancers. Mutation or dysregulation of the telomerase repression pathway may cause reactivation or upregulation of telomerase expression in cancer. Substances, identified in the methods, can be used to block transcription from the TR gene promoter through interaction of the 5' regulatory sequences. These substances, e.g. antisense oligonucleotides, transcription factors, peptide nucleic acids and factors that disrupt signal transduction, are useful for cancer therapy. In particular, gene therapy vectors
                                                             This invention describes a novel composition comprising human telomerase having at least 2000-fold (preferably at least 6000-fold) increased relative purity compared with crude extract of cells from adenovirus-transformed kidney cell line. The composition is useful for eliciting an immune response in animals and may therefore be used as a vaccine for treating diseases associated with the overexpression of telomerase e.g. cancer. AAZ23626-Z23637 represent oligonucleotides used in the isolation of human clone 28-1 which contains a fragment of the human telomerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mouse and human telomerase RNA gene promoters, useful for tumor specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Telomerase RNA; TR; promoter; cytotoxin; cancer; neoplasia; hTR; gene therapy; thymidine kinase gene; anticancer therapy; human; mutagenesis; PCR primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human telomerase RNA gene (hTR) promoter specific primer H11d.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 20; DB 1; Length 20;
Pred. No. 1.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 0 A; 9 C; 4 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
4.4%; Score 20; DB
Best Local Similarity 100.0%; Pred. No. 1.3
Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                          described in the method of the invention
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Disclosure; Col 49-50; 34pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Fig 12; 109pp; English
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The invention relates to promoter regions from mouse and human telomerase
RNA (TR) component genes. The TR gene promoter can be linked to a
Cancer, especially a gene encoding a cytotoxin, for therapy of
Cancer, especially neoplasias. The telomerase is necessary for the
Cancer, especially neoplasias. The telomerase is necessary for the
Cancer, especially neoplasias. The telomerase is necessary for the
Coursegulation of the telomerase repression pathway may cause reactivation
or upregulation of telomerase expression in cancer. Substances,
Cidentified in the methods, can be used to block transcription factors,
cancer through interaction of the 5' regulatory sequences. These
CS substances, e.g. antisense oligonuclectides, transcription factors,
c peptide nucleic acids and factors that disrupt signal transduction, are
Useful for cancer therapy. In particular, gene therapy vectors
(sepecially pGT62-codAupp) comprising the promoter and a viral thymidine
kinase gene can be used to convert a prodrug, e.g. gancyclovir, so that
c neoplasia can be controlled or treated. Direct down-regulation of
telomerase RNA gene through manipulation of transcription factors may be
effective anticancer therapy and the cloning of the hTR gene promoter
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(especially pGT62-codAupp) comprising the promoter and a viral thymidine kinase gene can be used to convert a prodrug, e.g. gancyclovir, so that neoplasia can be controlled or treated. Direct down-regulation of transcription of telomerase RNA gene through manipulation of transcription factors may be effective anticancer therapy and the cloning of the hTR gene promoter allows the analysis of therapeutic molecules which modulate hTR promoter activity. Sequences AA207096-321 represent PCR primers used in cloning
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mouse and human telomerase RNA gene promoters, useful for tumor specific
                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1) TR; promoter; cytotoxin; cancer; neoplasia; hTR;
thymidine kinase gene; anticancer therapy; human;
                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                                                           Score 20; DB 1; Length 20;
Pred. No. 1.3e+02;
                                                                                                                                                                                                                                                                  0; Indels
                                                                                                                                             and mutagenisis of human TR gene (hTR) promoter region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human telomerase RNA gene (hTR) specific primer hTRe.
                                                                                                                                                                                   Sequence 20 BP; 1 A; 4 C; 12 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                               4.4%; Scor.
100.0%; Pred. No. 1...
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure, Fig 6; 109pp, English.
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                                                                                                                                                                                                                                                                                                            17 GCCTGGGAGGGGTGGTGGCC 36
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                                                                                                                                                                                                                                                                      20; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1999-479183/40.
                                                                                                                                                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PCR primer; 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 164
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present sequence represents a peptide nucleic acid molecule which hybridises to the mRNA component of mammalian telomerase, and inhibits telomerase activity. Telomerase is a ribonucleoprotein enzyme that synthesizes one strand of the telomeric DNA, using as a template an 11 nucleotide sequence contained within the RNA component of the enzyme. The protectide sequence contained within the RNA component of the enzyme. The bases, which include the sequence GTTAGG. The uncharged nature of the PNA backbone increases the melting temperature of associating strands, increases the rast of association with targeted nucleic acids, and affords greater resistance of degradation by proteases or nucleases. The therapeutic PNAs may be used for treating disease conditions such as cancers, neoplasia, hyperplasia, neurodegenerative diseases, aging, human immunodeficiency virus (HTV) infection/AIDS (acquired immunodeficiency syndrome) and associated pathologies, fungal infections, and other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New peptide nucleic acid (PNA) compounds that inhibit telomerase activity in mammalian cells is useful as probes to detect the RNA component of a mammalian telomerase.
allows the analysis of therapeutic molecules which modulate hTR promoter activity. Sequences AA207623-80 represents PCR primers for amplifying human TR gene (hTR) promoter sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                         Peptide nucleic acid; PNA; telomerase; ribonucleoprotein enzyme; cancer; inhibitor; neoplasia; neurodegenerative disease; aging; hyperplasia; AIDS; HIV; fungal infection; forensic identification; detect; tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1. .20 a /*tag= a /*tag= bptide nucleic acid molecule, where N-(2-aminoethyl)glycine units are linked to nucleotide bases via glycine amino N through a methylenecarbonyl linker"
                                                                                                                                         Gape
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                                                                                                        Length 20;
                                                                                                                                       0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                           PNA sequence #41 used to inhibit telomerase activity.
                                                                   Sequence 20 BP; 3 A; 5 C; 8 G; 4 T; 0 U; 0 Other;
                                                                                                                        1.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Shay JW, Norton JC,
                                                                                                        DB 1;
                                                                                                               100.0%; Pred. ....
                                                                                                      Score 20;
Pred. No.
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                                                                                                                                                                            410 CTGAGCTGTGGGACGTGCAC 429
                                                                                                                                                                                                     CTGAGCTGTGGGACGTGCAC 20
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                                                                                                        4.48;
                                                                                                                                                                                                                                                                                                     AAA37583 standard; DNA; 20
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                                                                                                                     Best Local Similarity 100
Matches 20; Conservative
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misc_feature
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                                        AAA37583;
                                                                                                          Query Match
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ID AAA3
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diseases characterized by abnormal telomere metabolism or telomerase activity, in combination with antineoplastic and other cytotoxic or cytostatic agents, antifungal agents, and other nucleotides. PNAs may be used for molecular diagnostics, labelled PNAs are used as hybridization probes to detect or quantitate polynucleotides having a human telomerase RNA (hTR) sequence. PNA probes are also used for forensic identification of individuals, e.g. paternity testing, based on hTR gene restriction fragment length polymorphism (RFLP) pattern. PNAs are also useful as probes to detect the RNA component of a mammalian telomerase and as inhibitors of telomerase activity. The method of the present invention allows cancerous conditions to be detected with increased confidence and possibly at an earlier stage, before cells are detected as cancerous based on pathological characteristics. The diagnostic and prognostic methods of the present invention can be used to detect an immortal or neoplastic cell or tumour tissue or cancer of any origin, provided the cell expresses telomerase activity and its RNA component
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New peptide nucleic acids that hybridizes to the RNA component of mammalian telomerase, useful for treating or preventing cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     neurodegenerative disease; neoplasia; hyperplasia; HIV; AIDS;
human immunodeficiency virus; acquired immunodeficiency syndrome;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4.4%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 1.3e+02; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 3 A; 5 C; 5 G; 7 T; 0 U; 0 Other;
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AAS15454 standard; DNA; 20 BP.
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09-APR-1997;
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The present invention relates to peptide nucleic acids (PNAs), comprising a sequence of 6-25 nucleobases, that inhibit telomerase activity in mammalian cells by hybridising to the RNA component of mammalian cells by hybridising to the RNA component of mammalian cells by hybridising to the RNA component of telomerase and as inhibitors of telomerase activity, or to detect and/or quantitate polymucleotide having the human telomerase RNA component (hTR) sequence, as well as in forensic identification of individuals, such as paternity testing or identification of criminal cuspects or unknown descendants based on the hTR gene RFLP pattern. The PNA can be further used for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases. The PNAs in combination with other pharmacceuticals (such as antineoplastic or cytostatic agents) can be used for treating neoplast, hyperplasia, human immunodeficiency virus (HIV) infections, acquired immunodeficiency synchome (AIDS) and associated pathologies, and other diseases characterised by abnormal telomera metabolism or telomerase cutivity. The present sequence represents one of the PNA sequences of the
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breast; prostate; colon; mixed cell leukaemia; Hodgkin's disease;
fertility; inflammatory condition; tumour; cancer; veterinary;
immunosuppression; telomerase inhibitor; ss.
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inflammation, lymphoproliferative diseases, autoimmune disease, or
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/note= "N3'-P5' phosphoramidate linkages"
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                                                     English
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                  neurodegenerative diseases.
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                                                      29; 46pp;
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Best Local Similarity
                                                     Example 1; Col
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modified base
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Matches
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The invention relates to polynucleotide inhibitors (I) and methods for inhibiting telomerase activity. (I) are useful in inhibiting telomerase activity. (I) are useful in inhibiting telomerase activity and proliferation of a telomerase positive cell, and in manufacturing a medicament for inhibiting telomerase activity in a cell and in treating telomerase-mediated condition or disease, such as adenocarcinoma of breast, prostate or colon, mixed cell leukaemia, Hodgkin's disease, fertility and inflammatory conditions. (I) are also useful in treating a tumour or in manufacturing a medicament for the treatment of tumour. The polynucleotide inhibitors may also be used in diagnostic assays for detecting RNA or DNA. Inhibition of telomerase activity in cells in vivo is useful in prophlactic and therapeutic methods of treating cancer and other disorders involving inappropriate expression of telomerase, and in treating veterinary proliferative diseases. Inhibition of telomerase in haematopoietic stem cells is useful for immunosuppression and for selectively down-regulating specific branches of the immune system. The present sequence represents human the method of the immune system. The present sequence represents human the method of
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New polynucleotide useful for inhibiting telomerase activity in cells, for treating telomerase-mediated condition or disease, such as cancers, tumors, Hodgkin's disease, or inflammatory conditions.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Antisense oligonucleotide for human telomerase, Oligo 14ab #1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4.4%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 1.3e+02; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Seguence 20 BP; 3 A; 3 C; 3 G; 11 T; 0 U; 0 Other;
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                                                                                                                                        Claim 8; Page 36; 48pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     99US-00420056.
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97US-00833377.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Weinrich SL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US6261556-B1
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04-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           24-OCT-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAS09477;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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AASO9477/C
AASO9477/C
XX
AASO9477/C
XX
AASO9477/C
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AASO9477/C
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AASO9477/C
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The sequence represents an antisense oligonucleotide used in the purification of human telomerase. The invention relates to a purified human telomerase. The invention relates to a purified human telomerase core enzyme protein comprising 2000-fold increased than a crude extract of cells from adenovirus-transformed purity compared with a crude activity and a molecular weight of 200-2000 kidney cell line (293 cells) and when associated with telomerase RNA component has DNA polymerase activity and a molecular weight of 200-2000 kilo Daltons (KDa). The purified telomerase is useful for inducing a humoral or cell-mediated immune response in an animal. Purified telomerase or immunogenic fragments are useful as vaccines for treating diseases associated with over-expression of telomerase, such as cancer as affinity agents in insolating the proteins and for detecting the proteins are sell or tissue. Identification of telomerase aids in diagnosis of cancer or pre-cancerous states.

Telomerase and/or telomerase associated proteins are also useful for screening compounds to identify agents that alter the association of telomerase-association or EF2H with telomerase
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The sequence represents a biotinylated antisense oligonucleotide used in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Purified human telomerase, useful for inducing immune response in animals, comprises several thousand folds increased purity compared with cytoplasmic crude cell preparations.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pruzan RA;
                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Antisense oligonucleotide for human telomerase, Oligo 14ab #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human, Telomerase, vaccine, antibody, cancer; EF2H; nucleolin; antisense oligonucleotide; Oligo 14ab; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Weinrich SL, Atkinson EM, Lichtsteiner SP, Vasserot AP, Kealey \mathtt{JT}_{\mathsf{f}}
                                                                                                                                                                                                                                                                                                                                                      Length 20;
                                                                                                                                                                                                                                                                                                                                                                                       0; Indels
                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 0 A; 9 C; 4 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                      4.4%; Score 20; DB 1; Le
100.0%; Pred. No. 1.3e+02;
                                                                                                                                                                                                                                                                                                                                                                 100.0%; Preq. ...
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/mod_base= C
/note= "C is biotinylated"
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                                                                                                                                                                                                                                                                                                                                                                                                                             361 AGGCCGCAGGAAGAGGAACG 380
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1es 20; Conservative
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modified_base
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                                                                                                                                                                                                                                                                                                                                                        Query Match
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AAS09480/c
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US09436060A-16.rng.sl

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the purification of human telomerase. The invention relates to a purified human telomerase core enzyme protein comprising 2000-fold increased purity compared with a crude extract of cells from ademovinus-transformed kidney cell line (293 cells) and when associated with telomerase RNA component has DNA polymerase activity and a molecular weight of 200-2000 kilo Daltons (KDa). The purified telomerase is useful for inducing a humoral or cell-mediated immune response in an animal. Purified telomerase or immunogenic fragments are useful as vaccines for treating telomerase associated with over-expression of telomerase, which are useful as affinity agents in isolating the proteins and for detecting the presence of proteins in a sample, such as cell or tissue. Identification of telomerase and/or telomerase associated proteins are also useful for screening compounds to identify agents that alter the association of telomerase and/or telomerase associated proteins are also useful for screening compounds to identify agents that alter the association of telomerase associated proteins, such as nucleolin or EF2H with telomerase
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention provides a method for inhibiting or reducing the growth of a cell which involves administering to the cell a telomere damage inducing agent and a telomerase inhibitory agent. This can be used in the treatment of aberrant cell growth, including cancers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 growth of cell for treating cancer, comprising damage-inducing agent and telomerase inhibitory
                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                                                                                                                Score 20; DB 1; Length 20;
Pred. No. 1.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Telomerase; cancer; telomere damage; PCR primer; ss.
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                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 0 A; 9 C; 4 G; 7 T; 0 U; 0 Other;
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100.0%; Pred. No. ...
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                                                                                                                                                                                                                                                                                                                                                                                                                         361 AGGCCGCAGGAAGAACG 380
                                                                                                                                                                                                                                                                                                                                                                                                                                                      AGGCCGCAGGAAGAGGAACG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human telomerase PCR primer #2.
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nes 20; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Wientjes G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      agent to the cell.
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                                                                                                                                                                                                                                                                                                                                                     Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 170
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this represents an antisense oligonuclectide to the RNA component of human telomerase (hTR). The invention relates to an activator-antisense complex that comprises: (a) an antisense oligonuclectide, complementary to a 12-25 nuclectide portion of hTR, with a hydroxyl moiety at the first to a 12-25 nuclectide portion of hTR, with a hydroxyl moiety at the first card; and (b) a linker attached to the first end, and (c) an activator of RMsse L attached to the linker. The activator-antisense complex may be used for inhibiting the growth of a telomerase-expressing malignant cells or tumour. The complex is used to specifically cleave the ribonuclectide or tumour, renal cell cancer, mean and prostate cancer. Many other tumour, renal cell cancer, mean and prostate cancer. Many other canting and related disorders, may be treated including various acute and chronic leukemias, polychemia vera, Hodgkin's and non-Hodgkin's lymphomas, multiple myeloma, Waldenstrom's macroglobulinemia, Hodgkin's lymphomas and selate myeloma, waldenstrom's macroglobulinemia, carcinomas. The complex is preferably administered in combination with a chemotherapeutic agent, particularly either displatin, doxornation with a chemotherapeutic agent, particularly either displatin, doxornation with
                                                                                                                                                                                                                                                                                   Human; telomerase; hTR; activator-antisense complex; malignant; enzyme; cleave; brain; tumour malignant glioma; breast tumour; renal cell cancer; melanoma; prostete cancer; leukemia; polytchemia vera; myeloma; sarcoma; Hodgkin's lymphoma; Waldenstrom's macroglobulinemia; heavy chain disease; carcinoma; chemotherapeutic; antisense; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   mitomycin, daunorubicin, bleomycin, actinomycin D, or neocarzinostatin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New RNase L activator-telomerase antisense complex - useful to inhibit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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0
                                                                                                                                                                                                                                                  Human telomerase RNA (hTR) component antisense oligonucleotide.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      telomerase activity in telomerase-expressing malignancies
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Li G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cowell JK,
20
                                  1 GGGTTGCGGAGGGTGGGCCT 20
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98US-00018125.
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                                                                                                                                   AAV68462 standard; DNA; 19
                                                                                                                                                                                                                 (first entry)
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les 19; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
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03-FEB-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic
                                                                                                                                                                              AAV68462;
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Matches
                                                                                              RESULT 171
                                                                                                                     AAV68462/
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GTGCTTTTGCTCCCGGGC 94

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Gaps

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Query Match 4.4%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 1.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels

BP.

AAA37602 standard; RNA; 19

RESULT 173

AAA37602

AAA37602;

Telomerase target sequence.

15-AUG-2000 (first entry)

8

19 GTGCTTTTGCTCCCGCGC 1

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Sequences shown in AAV41169 to AAV41181 represent antisense
oligonucleotides to the RNA component of human telomerase (hTR). These
antisense oligonucleotides specifically hybridise to a nucleotide
sequence within an accessible region of the hTR, but that does not
hybridise to a sequence within the template region of hTR. These
component of human telomerase in a sample. This is useful for diagnosing
cancer (especially neuroblastoma, bladder, colon and prostate cancer),
and providing prognosis for a cancer patient. The inhibitory
oligonucleotides can inhibit the telomerase activity level in a cell by
interfering with transcription of the RNA component, decreasing the half-
life of the telomerase RNA component transcript, inhibiting the
polymerase activity of telomerase holeenzyme, or inhibiting the
polymerase activity of telomerase activity in both cultured cells and in
cells in vivo. They can be used in therapeutics for treating or
munosuppression, and for selectively down-regulating specific branches
immunosuppression, and for selectively down-regulating specific branches
of the immune system.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New polynucleotide(s) anti:sense to human telomerase - used for detecting or inhibiting human telomerase, e.g. for treating cancers, contraception, immuno-suppression or treating infection.
                                                                                                                                                RNA component; human telomerase; antisense oligonucleotide; infection; neuroblastoma; bladder cancer; colon cancer; prostate cancer; cancer; contraception; sterilisation; immunosuppression; therapeutic; hTR; immune system down-regulation; anti-inflammatory therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           inflammatory therapies or for treating infections by, e.g. yeast, parasites or fungi
                                                                                                            RNA component of human telomerase (hTR) antisense oligo 21ab.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4.2%; Score 19; DB 1; Length 19; 00.0%; Pred. No. 1.5e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 11; Page 65; 80pp; English
BP
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AAV41176 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Kim NW, Wu F, Kealey JT,
                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1998-377670/32.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (GERO-) GERON CORP.
                                                                                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                        WO9828442-A1
                                                                                                                                                                                                                                                                                                                                                                                 19-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                     20-DEC-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                        20-DEC-1996;
                                                                         08-OCT-1998
                                                                                                                                                                                                                                                                                                                                            02-JUL-1998.
                                                                                                                                                                                                                                               Synthetic.
                                   AAV41176;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
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The present sequence represents a telomerase target sequence. The invention relates to an antisense oligonuclectide used as a control sequence alongside a peptide nucleic acid molecule which hybridises to the mammalian telomerase, and inhibits telomerase activity. Telomerase is a ribonucleoprotein enzyme that synthesizes one activity. Telomerase is a ribonucleoprotein enzyme that synthesizes one strand of the telomeric DNA, using as a template an il nucleotide contained within the RNA component of the enzyme. The invention contained within the RNA component of the enzyme. The invention relates to PNA molecules having a sequence of row more than 25 bases, which include the sequence GTTARGG. The uncharged nature of the PNA backbone increases the melting temperature of associating strands, conficience of association with targeted nucleic acids, and the increases the rate of association with targeted nucleic acids, and concrets, neoplasia, hyperplasia, neurodegenerative diseases or nucleases. The charapeutic PNAs may be used for treating disease conditions such as cancers, neoplasia, hyperplasia, neurodegenerative diseases, aging, human immunodeficiency virus (HIV) infection/AIDS (acquired immunodeficiency syndrome) and associated pathologies, fungal infections, and other cytocoxic circ diseases characterized by abnormal telomerase activity, in combination with antineoplastic and other cytocoxic cytostatic agents, antifungal agents, and other nucleotides. PNAs may be used for molecular diagnostics, labelled PNAs are used as hybridization cytoscatic agents, antifungal agents, and other nucleotides. PNAs may be used for molecular diagnostics, labelled PNAs are used as hybridization of individuals, e.g. paternity testing, based on hTR gene restriction for probes to detect the RNA component of a mammalian telomerase and as probes to detect the RNA component of a mammalian telomerase and allows cancerous conditions to be detected with increased confidence and New peptide nucleic acid (PNA) compounds that inhibit telomerase activity in mammalian cells is useful as probes to detect the RNA component of a Peptide nucleic acid; PNA; telomerase; ribonucleoprotein enzyme; cancer; possibly at an earlier stage, before cells are detected as cancerous based on pathological characteristics. The diagnostic and prognostic methods of the present invention can be used to detect an immortal or neoplastic cell or tumour tissue or cancer of any origin, provided the inhibitor, neoplasia, neurodegenerative disease, aging, hyperplasia, AIDS, HIV, fungal infection, forensic identification, detect, tumour, paternity testing; ss. Corey DR; expresses telomerase activity and its RNA component Norton JC, Shay JW, Example 2; Col 37; 45pp; English. 97US-00838545. 96US-00630019 Piatyszek MA, (TEXA) UNIV TEXAS SYSTEM mammalian telomerase. WPI; 2000-292432/25. 09-APR-1997; 09-APR-1996; 04-APR-2000. US6046307-A. Wright WE, Synthetic.

Sequence 19 BP; 7 A; 5 C; 3 G; 0 T; 4 U; 0 Other;

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Gaps

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0; Indels

100.0%; Prec.

19; Conservative

Matches

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Best Local Similarity

148 CCACCGTTCATTCTAGAGC 166

19 ccaccerrcarrcragaec 1

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The present sequence represents an antisense polyamide nucleic acid (PNA)), which is covalently linked to a RNase Lactivator moiety. The oligonuclectide is used in the method of the activator moiety. The oligonuclectide is used in the method of the invention. The specification describes a method of cleaving a specific RNA, optionally contained within a cell. The method comprises hybridising the RNA with a chimeric molecule of antisense PNA and RNAse L activator molecule is useful for treating Respiratory Syncytial Virus infection, inhibiting tumour growth, treating viral infection (e.g. human immunodeficiency virus (HIV)), autosomal dominant diseases caused by a mutant gene, chronic myelogenous leukemias, and Crohns disease and other inflammatory conditions. It is also useful for sequence-specific inhibition of a targeted mRNA in order to effectively knock out the associated gene as an aid to elucidating the function of the encoded
                                   ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cleaving a specific RNA comprising hybridizing the RNA with a chimeric molecule consisting of antisense polyamide nucleic acid (pna) and RNAse L activator moieties, useful for treating e.g. cancer and inflammatory
                                                                                                                                                                                                                                                                                                                                    Antisense polyamide nucleic acid; peptide nucleic acid; PNA; RNase L activator; RNA cleavage; antisense PNA; tumour growth; respiratory syncytial virus infection; viral infection; human immunodeficiency virus; HIV; autosomal dominant disease; chronic myelogenous leukemia; Crohns disease; inflammatory condition; ss.
                                   Gaps
                                   .
0
                                                                                                                                                                                                                                                                                                    Nucleotide sequence of an antisense polyamide oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Van Der Marel GA;
Length 19;
                                   0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 19 BP; 6 A; 5 C; 8 G; 0 T; 0 U; 0 Other;
Score 19; DB 1; Pred. No. 1.5e+02; 4; Mismatches 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Verheijen JC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (USSH ) US DEPT HEALTH & HUMAN SERVICES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 25; Page 32; 41pp; English.
                                                                        62
                                                                                                         BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          98US-0099173P.
                                                                          GTCTAACCCTAACTGAGAA
                                                                                                                                                                                                  AAZ99775 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Van Boom JH,
                                                                                                                                                                                                                                                                     (first entry)
   Query Match
Best Local Similarity 78.9
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (UYLE-) RIJKSUNIV LEIDEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2000-256971/22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200014219-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         02-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            04-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Torrence PF,
                                                                                                                                                                                                                                                                       12-JUL-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16-MAR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
                                                                                                                                                                                                                                    AAZ99775;
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                                                                                                                                                               RESULT 174
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The present invention relates to peptide nucleic acids (PNAs), comprising a sequence of 6-25 nucleobases, that inhibit telomerase activity in nammalian cells by Mybridising to the RNA component of mammalian cells by Mybridising to the RNA component of mammalian telomerase and as inhibitors of telomerase activity, or to detect and/or quantitate polynucleotide having the human telomerase RNA component (HTR) sequence, as well as in forensic identification of individuals, such as paternity testing or identification of criminal suspects or unknown descendants based on the MTR gene RFLP pattern. The PNAs can be further used for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases, and combination with other pharmaceuticals (such as antineoplastic or cytostatic agents) can be used for treating neoplasia, hyman immunodeficiency virus (HIV) infections, acquired immunodeficiency syndrome (AIDS) and associated pathologies, and other diseases characterised by abnormal telomere metabolism or telomerase to activity. The present sequence repersenting hTR RNA strand #3 is used to test the inhibition of telomerase activity by the PNAs of the present
                                                                                                                                                                                                                                                                Mammalian; forensic; paternity testing; human telomerase RNA component; hTR gene RFLP pattern; cancer; inflammation; lymphoproliferative disease; autoimmune disease; neurodegenerative disease; neoplasia; hyperplasia; HIV; AlDS; human immunodeficiency virus; telomere metabolism; cytostatic; acquired immunodeficiency syndrome; anti-inflammatory; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New peptide nucleic acids that hybridizes to the RNA component of mammalian telomerase, useful for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 19; DB 1; Length 19;
Pred. No. 1.5e+02;
4; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 19 BP; 7 A; 5 C; 3 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Piatyszek MA, Corey DR,
                                                                                                                                                                                                                               Human telomerase RNA (hTR) strand #3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 2; Col 37-38; 46pp; English.
                  19 Grecririscreceded 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 Similarity 78.9%;
15; Conservative
neurodegenerative diseases
                                                                                                                AAS15473 standard; RNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (TEXA ) UNIV TEXAS SYSTEM.
                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Wright WE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2001-638024/73
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            09-APR-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                             US6294650-B1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      08-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                09-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-SEP-2001.
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                                                                                                                                                         AAS15473;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Shay JW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Loca
Matches
                                                                            RESULT 175
                                                                                                  AAS15473
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1 GUCUAACCCUAACUGAGAA 19 GTCTAACCCTAACTGAGAA 62

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44

8 g

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Gaps

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4.2%; Score 19; DB 1; Length 19; 100.0%; Pred. No. 1.5e+02; ive 0; Mismatches 0; Indels

Query Match Best Local Similarity 100.0 Matches 19; Conservative

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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA characteries. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprises esnes and antisense of single stranded. They further comprises sense and antisense of single stranded. They further comprises enses and antisense of single stranded. They further comprises and antisense of single stranded. The single sinclude short interfering RNA (shRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (shRNA). The siNAs can be unmodified or chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA, conjugates and/or complexes of siNA; and vectors that express sink. The siNAs are used to modulate expression of the telomerase genes in calls, tissue candour complexes of siNA; and vectors there express sink. The siNAs are used to modulate expression of the telomerase genes in calls, tissue captaints for the treatment of a variety of conditions. They may be used for treatming cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases. Ce sy multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying sene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the upper strand of a human TERC-targeted double-stranded sink, which is identical to the c-fos transcript target sequence.
                                                                                                                                                                                                                      Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthitic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short haltpin RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
                                                                                                                                                                                Human TERC transcript target sequence/siNA upper strand, SEQ ID 20.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 3; SEQ ID NO 20; 145pp; English.
                                             ADF93303 standard; RNA; 19 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2002US-0358580P
2002US-0363124P
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2002US-0396600P.
2002US-0406784P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mcswiggen J, Beigelman L;
                                                                                                                                     (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO2003070742-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       06-JUN-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20-FEB-2002;
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                                                                                                                                     26-FEB-2004
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                                                                                          ADF93303;
RESULT 176
                       ADF93303
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                                                                                                                                                                                                                                                                                                                                                                                                                         Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthitic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; sinA; short interfering RNA; albNA; double-stranded RNA; micro-RNA; miRNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                        Human TERC transcript target sequence/siNA upper strand, SEQ ID 24.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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                                                Query Match 4.2%; Score 19; DB 1; Length 19; Best Local Similarity 89.5%; Pred. No. 1.5e+02; Matches 17; Conservative 2; Mismatches 0; Indels
                G; 0 T; 2 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 3; SEQ ID NO 24; 145pp; English.
                                                                                                                                  248 CTGGAGGCCGCGGTCGGCC 266
                                                                                                                                                                         1 cuddaddccccccccuceccc 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2002US-0358580P.
2002US-0363124P.
2002US-0386782P.
2002US-0406784P.
2002US-0406378P.
2002US-0409293P.
                                                                                                                                                                                                                                                                      ВР
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                Sequence 19 BP; 1 A; 7 C; 9
                                                                                                                                                                                                                                                                      ADF93307 standard; RNA; 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mcswiggen J, Beigelman L;
                                                                                                                                                                                                                                                                                                                                                  26-FEB-2004 (first entry)
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17-JUL-2002;
29-AUG-2002;
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                                                                                                                                                                                                                                  RESULT 177
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   used for treating cancer, restenois, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siMas are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene represents the upper strand of a human TERC-targeted double-stranded sIMA, which is identical to the c-fos transcript target sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense
                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human TERC transcript target sequence/siNA upper strand, SEQ ID 29.
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 transplants for the treatment of a variety of conditions.
                                                                                                                                                                                                                                        4.2%; Score 19; DB 1; Length 19; 84.2%; Pred. No. 1.5e+02; ive 3; Mismatches 0; Indels
                                                                                                                                                                                                   Sequence 19 BP; 0 A; 7 C; 9 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 3; SEQ ID NO 29; 145pp; English.
                                                                                                                                                                                                                                                                                                               320 CCGCGGGTCTCTCGGGGGC 338
                                                                                                                                                                                                                                                                                                                                    1 CCGCGGGUCUCUCGGGGGC 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 11-MAR-2002; 2002US-0363124P.
06-JUN-2002; 2002US-0363124P.
17-JUL-2002; 2002US-039660P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-040878P.
                                                                                                                                                                                                                                                                                                                                                                                                                                            ADF93312 standard; RNA; 19 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
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                                                                                                                                                                                                                                                         Local Similarity 84.2
les 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADF93312;
                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                           Best Loca
Matches
                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 178
888888888888888888
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cc regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short in the faring RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised. Serpressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, c.g. multiple sclerosis, lupus erythmatosus, AlDS, macular degeneration, stin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence was which is adaptically the content of a human TERC targeted double-stranded
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                      siNA, which is identical to the c-fos transcript target sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4.2%; Score 19; DB 1; Length 19; 78.9%; Pred. No. 1.5e+02; trive 4; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 19 BP; 3 A; 4 C; 8 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human TERC siNA lower strand, SEQ ID 287.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       410 CTGAGCTGTGGGACGTGCA 428
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 CUGAGCUGUGGGACGUGCA 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          , 2002US-0358580P.
; 2002US-0363124P.
; 2002US-0386782P.
; 2002US-0396600P.
; 2002US-0406784P.
; 2002US-0408378P.
; 2003US-040833P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity 78.9
les 15; Conservative
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17-JUL-2002; 2
29-AUG-2002; 2
05-SEP-2002; 2
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WPI; 2003-689777/65.

29-AUG-2002; 2002US-0406784P. 05-SBP-2002; 2002US-0408378P. 09-SEP-2002; 2002US-0409293P. 15-JAN-2003; 2003US-0440129P.

(RIBO-) RIBOZYME PHARM INC.

Beigelman L;

WPI; 2003-689777/65. Mcswiggen J,

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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA connergulate expression of the one or more telomerase genes by RNA connergulate expression of mean telementatively are assembled from a sense oligonucleotide and an interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short interfering RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, and/or camplexes of siNA; and vectors that express in calls, tissue contained to modulate expression of the telomerase genes in calls, tissue capplants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, skin ulcers and rheumatoid arthritis. The sinas are also useful for drug screening, diagnosis, lupus erythematosus, ALDS, macular degeneration, skin ulcers and rheumatoid arthritis. The sinas are also useful for drug careening, diagnosis, lupus erythematosus, ALDS, macular degeneration, capping (e.g., of single nucleotide polymorphisms). The present sequence construction, and gene construction, and gene construction, and gene construction, and gene construction and such construction 
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                    New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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100.0%; Pred. No. 1.5e+02;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 19 BP; 5 A; 7 C; 4 G; 0 T; 3 U; 0 Other;
                                                                                       Example 3; SEQ ID NO 287; 145pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 302 AGAGTIGGCCTCTGTCAGC 320
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Matches 19; Conserv
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The invention relates to short interfering nucleic acids (sinA) which downregulate expression of the one or more telomerase genes by RNA interference. The sinNa may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and antiense regions, or alternatively are assembled from a sense oligonuclectide and an antisense oligonuclectide. Specifically, the sinNa include short hairpin RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs can be unmodified or chemically synthesised, can contain deoxyribonuclectides, and can be chemically synthesised, can contain deoxyribonuclectides, and can be chemically synthesised. Cexpressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be

New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.

Example 3; SEQ ID NO 291; 145pp; English.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, suddying gene function, and gene mapping (e.g., of single nuclecide polymorphisms). The present sequence represents the lower strand of a human TERC-targeted double-stranded
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 19 BP; 1 A; 7 C; 6 G; 0 T; 5 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
4.2%; Score 19; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 1.5e+02;
Matches 19; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    374 AGGAACGGAGCGAGTCCCC 392
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Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short historing RNA; siRNA; couble-stranded RNA; micro-RNA; miRNA; double-stranded RNA; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.

11-MAR-2002; 2002US-0363124P. 06-JUN-2002; 2002US-0386782P. 17-JUL-2002; 2002US-0396600P.

11-FEB-2003; 2003WO-US004088

20-FEB-2002;

WO2003070742-A1 Homo sapiens.

28-AUG-2003

Human TERC siNA lower strand, SEQ ID 291.

26-FEB-2004 (first entry)

ADF93564;

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neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic;
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Matches
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                                                                                                                                                                                                                                                                                                                                                                   The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and an intisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short harpin RNA (shRNA). The siNAs can be unmodified or chemically modified, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and vector or express sion of the telomerase genes in cells, tissue used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and treatment of a variety of conditions. They may be used for treating cancer, restenoisis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, c. g. multiple sclerosis, lubus erythematosus, ADDS, macular degeneration, contening, diagnosis, therappeutic target identification and validation, or surface the sinks are also useful for drug screening, diagnosis, therappeutic target identification and validation,
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mapping (e.g., of single nucleotide polymorphisms). The present sequence
represents the upper strand of a human TERC-targeted double-stranded
siNA, which is identical to the c-fos transcript target sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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                                                                                                                                                                                                                                                                                                     short interfering nucleic acid downregulates expression of the
                                                                                                                                                                                                                                                                                                                    telomerase gene useful e.g. for treatment and diagnosis of cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 19 BP; 0 A; 8 C; 4 G; 0 T; 7 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                              Example 3; SEQ ID NO 11; 145pp; English.
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                                                                                  20-FEB-2002; 2002US-0358580P.

11-MAR-2002; 2002US-0353124P.

06-JUN-2002; 2002US-0386782P.

17-JUL-2002; 2002US-0336600P.

29-AUG-2002; 2002US-0406784P.

05-SEP-2002; 2002US-0408378P.

09-SEP-2002; 2002US-040929P.

15-JAN-2003; 2003US-0440129P.
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                                                          11-FEB-2003; 2003WO-US004088
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Best Local Similarity 63.2'
                                                                                                                                                                                                                                                                          WPI; 2003-689777/65.
  WO2003070742-A1
                                                                                                                                                                                                                                               Mcswiggen J,
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                             28-AUG-2003
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₹8825X6
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA communications and antisense of interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense coligonacleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short interfering RNA (siRNA). The siNAs can be unmodified or chemically synthesised, can contain deoxyribonucleotides, and can be enemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complaces of siNA; and vectors that express siNAs are cused to modulate expression of the telomerase genes in cells, tissue capiants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically, protozoal), transplant rejection, or autoimmune or age-related diseases, ce.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug correcting, dharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence cepture, expressents the upper strand of a human TERC-targeted double-stranded crepresents the upper strand of a transcript target sequence.
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antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-granded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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Pred. No. 1.5e+02;
5; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 3; SEQ ID NO 16; 145pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1:||||:||||:
1 AUGUCAGCUGCUGGCCCGU 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2002US-0363124P.
2002US-0386782P.
2002US-0396600P.
2002US-0406784P.
2002US-0409293P.
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es 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (RIBO-) RIBOZYME PHARM INC.
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                                                                                                                                                                                                                                                                                                                                                   WO2003070742-A1.
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05-SEP-2002;
09-SEP-2002;
                                                                                                                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      11-MAR-2002;
06-JUN-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20-FEB-2002;
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Length 19;

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Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
                                                                    Human TERC transcript target sequence/siNA upper strand, SEQ ID 17.
                                                                                                                                                                                                                                                                      2002US-0363124P.
2002US-0386782P.
2002US-0396600P.
2002US-0406784P.
        ADF93300 standard; RNA; 19 BP
                                                                                                                                                                                                                                                                                                              05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409293P.
15-JAN-2003; 2003US-0440129P.
                                                                                                                                                                                                                                          11-FEB-2003; 2003WO-US004088
                                                                                                                                                                                                                                                                                                                                                      (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                          Beigelman L;
                                                (first entry)
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                                                                                                                                                                                                  WO2003070742-A1.
                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                       11-MAR-2002;
06-JUN-2002;
                                                                                                                                                                                                                                                                                                    29-AUG-2002;
                                                                                                                                                                                                                                                                                                                                                                          Mcswiggen J,
                                                                                                                                                                                                                                                                                           17-JUL-2002;
                                                26-FEB-2004
                                                                                                                                                                                                                      28-AUG-2003
                            ADF93300;
ADF93300
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and matisense regions, or alternatively are assembled from a sense oligonuclectide and an antisense oligonuclectide. Specifically, the siNAs include short natisense oligonuclectide. Specifically, the siNAs include short hairpin RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonuclectides, and can be chemically synthesised, can contain deoxyribonuclectides, and can be chemically synthesised, captace to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and cransplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration skin ulcers and rheumatoid arthritis. The sINAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the upper strand of a human TERC-targeted double-stranded New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer. siNA, which is identical to the c-fos transcript target sequence. Example 3; SEQ ID NO 17; 145pp; English.

Seguence 19 BP; 1 A; 10 C; 5 G; 0 T; 3 U; 0 Other;

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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antiense regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short hairpin RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs can be unmodified or chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised. Service or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA, conjugates and/or complexes of siNA, and vectors that express sine. The siNAs are used to modulate expression of the telomerase genes in cells, tissue candout the treatment of a variety of conditions. They may be transplants for the treatment of a variety of conditions. They may be used to modulate concerned the sine of the telomerase genes in cells. They may be used to modulate concerned the sine of the conditions diseases (specifically concerned the sine of the telomerase genes in cells. They may be used to modulate concerned the sine of the sine of the telomerase seases (specifically concerned the sine of the sine of the telomerase seases (specifically concerned the sine of sine of the sine of sine 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    neuroprotective, anti-HIT, ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; sinA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
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                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                              Human TERC transcript target sequence/siNA upper strand, SEQ ID 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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0
Score 19; DB 1; Length 19;
Pred. No. 1.5e+02;
3; Mismatches 0; Indels
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2002US-0386782P.
2002US-039660P.
2002US-0406784P.
2002US-0408378P.
                                                                                                                                                                                                                                                                                      ВР
                                                                                                      194 TTCGCCCTCCCGGGGACC
  4.2%;
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                                                                                                                                                                                                                                                                                      ADF93306 standard; RNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (RIBO-) RIBOZYME PHARM INC.
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                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2003-689777/65.
  Query Match
Best Local Similarity
Matches 16; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20-FEB-2002;
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e.g. multiple sclerosis, lupus erythemarobus, hard, margine useful for drug skin ulcers and rheumatoid arthritis. The siMas are also useful for drug screening, diagnosis, therapeutic taget identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the upper strand of a human TBRC-targeted double-stranded siNA, which is identical to the c-fos transcript target sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiartitic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering NNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hinterfering syresion modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human TERC transcript target sequence/siNA upper strand, SEQ ID 31,
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Pred. No. 1.5e+02;
                                                                                                                                                                                                                           0; Indels
                                                                                                                                                 Sequence 19 BP; 3 A; 4 C; 7 G; 0 T; 5 U; 0 Other;
                                                                                                                                                                                                                           5; Mismatches
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                                                                                                                                                                                                                                                                302 AGAGTIGGGCTCTGTCAGC 320
                                                                                                                                                                                                                                                                                      AGAGUUGGCUCUGUCAGC 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20-FEB-2002; 2002US-0358580P.
11-MAK-2002; 2002US-0353124P.
06-JUN-2002; 2002US-0386782P.
17-JUL-2002; 2002US-0396600P.
9-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-040873PP.
                                                                                                                                                                                                                                                                                                                                                                                                ADF93314 standard; RNA; 19 BP
                                                                                                                                                                                  4.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                             14; Conservative
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                                                                                                                                                                                                           Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          26-FEB-2004
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                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                             RESULT 185
                                                                                                                                                                                                                             Matches
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ean contain decoxyribonuclectides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vivo of elivery of siNA, conjugates and/or complexes of siNA, and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explaints or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, ALDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the upper strand of a human TERC-targeted double-stranded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; anti-HIV; ophthalmological; elomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
hairpin RNA (shRNA). The sinAs can be unmodified or chemically modified,
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                                                                                                                                                                                                                                                                                                                                    siNA, which is identical to the c-fos transcript target sequence
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                                                                                                                                                                                                                                                                                                                                                                                                               4.2%; Score 19; DB 1; Length 19; 84.2%; Pred. No. 1.5e+02;
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                                                                                                                                                                                                                                                                                                                                                                      Sequence 19 BP; 5 A; 7 C; 4 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        3; Mismatches
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2002US-0386782P.
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2002US-0406784P.
2002US-0408378P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity 84.2
Matches 16; Conservative
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11-MAR-2002; 2
06-JUN-2002; 2
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29-AUG-2002;
05-SEP-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 186
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short interfering RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised. Expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue cyplants for the treatment of a variety of conditions. They may be used for treathing cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence mapping (e.g., of single nucleotide polymorphisms). The present sequence
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Pred. No. 1.5e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 4.2%; Score 19; UB 1
Best Local Similarity 100.0%; Pred. No. 1.5e
                                 Example 3; SEQ ID NO 290; 145pp; English.
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2002US-0386782P.
2002US-0396600P.
2002US-0406784P.
2002US-0408378P.
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05-SEP-2002;
09-SEP-2002;
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06-JUN-2002;
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Gaps

WO2003070742-A1 Homo sapiens

28-AUG-2003.

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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonuclectide and an interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonuclectides, and can be chemically synthesised. Can contain deoxyribonuclectides, and can be chemically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA: conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue cused to modulate expression of the telomerase genes in cells, tissue cused to modulate expression of the telomerase genes in cells, tesque cused transplants for the treatment of a variety of conditions. They may be used for transplants for the treatment of a variety of conditions. They may be used for transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug correining, diagnosis, therapelutic target identification and and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence correspond to a human TRRC-targeted double-stranded diseases correspond to a human transporter that part and correspond to a human transporter to the present engagement of a human transferior that presents the upper strand of a human transferior that present strand of a human transferior that presents the upper strand of a human transferior that presents the upper strand of a human transferior that proverser and the upper strand of a human transferior the present engineering the present the proverser and the up
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                                                                                                                                                                                                           New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       siNA, which is identical to the c-fos transcript target sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4.2%; Score 19; DB 1; Length 19;
84.2%; Pred. No. 1.5e+02;
ve 3; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 19 BP; 1 A; 7 C; 8 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human TERC siNA lower strand, SEQ ID 276.
                                                                                                                                                                                                                                                                                            Example 3; SEQ ID NO 21; 145pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         266 CCGGGGCTTCTCCGGAGGC 284
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15-JAN-2003; 2003US-0440129P.
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                                                  (RIBO-) RIBOZYME PHARM INC.
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                                                                                                       Mcswiggen J, Beigelman L;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 84.2
Matches 16; Conservative
                                                                                                                                                            WPI; 2003-689777/65.
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and an trisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short interfering RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyriboucleotides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and vectors that express sinA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically contains) transplant rejection, or autoimmune or age-related diseases, e.g. e.g. multiple sclerosis, lupus erythematosus, ADS, mecular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the lower strand of a human TERC-targeted double-stranded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                     New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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100.0%; Pred. No. 1.5e+02;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 19 BP; 5 A; 6 C; 6 G; 0 T; 2 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human TERC siNA lower strand, SEQ ID 294.
                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 3; SEQ ID NO 276; 145pp; English
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                                                                                               2002US-0386782P.
2002US-0396600P.
2002US-0406784P.
                                                                                                                                                               2002US-0408378P
                                                                                                                                                                                                        15-JAN-2003; 2003US-0440129P
                                                                                                                                                                                    2002US-0409293P
                 11-FEB-2003; 2003WO-US004088
                                                                                2002US-0363124P
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                                                                                                                                                                                                                                                                                           Mcswiggen J, Beigelman L;
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05-SEP-2002;
09-SEP-2002;
                                                                                                  06-JUN-2002;
                                                                                                                       17-JUL-2002;
                                                          20-FEB-2002;
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The invention relates to short interfering nucleic acids (siNA) which commission of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and antisense cregions, or alternatively are assembled from a sense oligonuclectide and antisense oligonuclectide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (shRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonuclectides, and can be chemically synthesised, can contain deoxyribonuclectides, and can be chemically synthesised, can complexes of siNA; and vectors that express in Rni invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the traatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically process), intenting are selected, in the sinas are also useful for drug screening, diagnosis, lupus erythematosus, AIDS, macular degeneration, egenetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence mapping (e.g., of single nucleotide polymorphisms). The present sequence the single of a human TERC-targeted double-stranded
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                         short hairpin RNA; shRNA; expression modulation; gene therapy;
drug screening; diagnosis; therapeutic target identification;
pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
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short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA;
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100.0%; Pred. No. 1.5e+02;
.ive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                    20-FEB-2002; 2002US-0358580P.

11-WAR-2002; 2002US-0361124P.

05-UUN-2002; 2002US-0386782P.

17-UUL-2002; 2002US-0396600P.

29-AUG-2002; 2002US-0406784P.

05-SEP-2002; 2002US-0408378P.

09-SEP-2002; 2002US-040923P.

15-JAN-2003; 2003US-0440129P.
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ID ADF93301 standard; RNA; 19 BP.
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mcswiggen J, Beigelman L;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-689777/65.
                                                                                                                                                         WO2003070742-A1
                                                                                                                  Homo sapiens.
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNas may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense colligonacleotide. Specifically, the siNas include short interfering RNA (shRNA), double-stranded RNA, micro-RNA (miRNA) and short interfering RNA (shRNA). The siNas can be unmodified or chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, captessed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express sinA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and cransplants for the treatment of a variety of conditions. They may be used for treatming cancer, restenosis, induse explantence or age-related diseases, conditions and rheumatoid arthritis. The siNAs are also useful for drug stin ulcers and rheumatoid arthritis. The siNAs are also useful for drug serienning, diagnosis, therapeuter target identification and validation, serening, diagnosis, therapeute, expending gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the upper strand of a human TERC-target educate.
                                                                                                                      Cytostatic; vasotropic; protozoacide; immunosupprassive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
                                                                                   Human TERC transcript target sequence/siNA upper strand, SEQ ID 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     29-AUG-2002; 2002US-0406784P-
05-SEP-2002; 2002US-0408378P-
09-SEP-2002; 2002US-0490293P-
15-JAN-2003; 2003US-0440129P-
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2002US-0386782P.
2002US-0396600P.
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                                         (first entry)
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                                                                                                                                                                                                                                                                                                                                        Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         11-MAR-2002;
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                                         26-FEB-2004
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  ADF93301;
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The invention relates to short interfering nucleic acids (sINA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (shRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue candor complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases.

Scorp multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation,
                                                                                                                                                                                                                                                                                                           Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering uncleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        telomerase gene useful e.g. for treatment and diagnosis of cancer.
                                                                                                                                                                                                                                                                        Human TERC siNA lower strand, SEQ ID 280.
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212 CTGCGGCGGGTCGCCTGCC 230
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2002US-0386782P.
2002US-0396600P.
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2002US-0408378P.
2002US-0409293P.
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                                                                                                                                               ADF93553 standard; RNA; 19
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                                                                                                                                                                                                                                 26-FEB-2004 (first entry)
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06-JUN-2002;
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Gaps

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0; Indels

4.2%; Score 19; DB 1; Length 19; 84.2%; Pred, No. 1.5e+02;

3; Mismatches

16; Conservative

Best Local Similarity

Query Match Matches

Sequence 19 BP; 0 A; 8 C; 8 G; 0 T; 3 U; 0 Other;

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ADF93305;
                                                                                                                                                                                                                                                                                                                     RESULT 193
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (shRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also
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genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the lower strand of a human TBRC-targeted double-stranded
                                                                                                             Gaps
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                                                                                   4.2%; Score 19; DB 1; Length 19;
100.0%; Pred. No. 1.5e+02;
ive 0; Mismatches 0; Indels
                                                            Sequence 19 BP; 5 A; 6 C; 6 G; 0 T; 2 U; 0 Other;
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                                                                                                                                    176 ATGTCAGCTGCTGGCCCGT 194
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2002US-0363124P.
2002US-0386782P.
2002US-0396600P.
2002US-0406784P.
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                                                                                                                                                                                                                        ADF93568 standard; RNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (RIBO-) RIBOZYME PHARM INC.
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                                                                                                                                                                                                                                                                        (first entry)
                                                                              Query Match
Best Local Similarity 100.0
Matches 19; Conservative
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06-JUN-2002; 2
17-JUL-2002; 2
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                                                                                                                                                                                                   RESULT 192
                                         BiNA.
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relates to kits for the in vitro or in vivo delivery of siNA, conjugates and/or complexes of siNA, and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lugus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleoide polymorphisms). The present sequence represents the lower strand of a human TERC-targeted double-stranded
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100.0%; Pred. No. 1.5e+02;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 19 BP; 3 A; 4 C; 7 G; 0 T; 5 U; 0 Other;
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2002US-0386782P.
2002US-039660DP.
2002US-0406784P.
2002US-040823P.
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Best Local Similarity
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06-JUN-2002;
17-JUL-2002;
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05-SEP-2002;
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Mcswiggen J,

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CC The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA controlled and antisense obtained stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short can contain deoxyribonucleotides, and can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, can expressed from a vector or enzymatically synthesised. The invention also crelates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express sincells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically protocoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The sinAs are also useful for drug screening, disagnosis, lupus erythematosus, AIDS, macular degeneration, can empining (e.g., of single nucleotide polymorphisms). The present sequence mapping (e.g., of single nucleotide polymorphisms). The present sequence contains the sinal sector of a human tracet organisms of the paper strand of a human tracet organisms.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              siNA, which is identical to the c-fos transcript target sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4.2%; Score 19; DB 1; Length 19; 94.7%; Pred. No. 1.5e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1; Mismatches
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2002US-0408378P.
2002US-0409293P.
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2002US-0386782P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity 94.7
nes 18; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20-FEB-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-SEP-2002;
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and antiense regions, or alternatively are assembled from a sense oligonuclectide and an artisense oligonuclectide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (shRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonuclectides, and can be chemically synthesised, can contain deoxyribonuclectides, and can be chemically synthesised.

C expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthitis. The siMs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the lower strand of a human TERC-targeted double-stranded
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                 New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 19 BP; 7 A; 6 C; 5 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                        Example 3; SEQ ID NO 274; 145pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human TERC siNA lower strand, SEQ ID 275.
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                                                                      WPI; 2003-689777/65.
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Matches
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pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
                                         Homo sapiens.
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06-JUN-2002;
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                                                                                                            28-AUG-2003
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                                                                                                                                                                                                                                                                                                                                                         The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense cregions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include and short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short can contain deoxyribonucleotides, and can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, and or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically proteozal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AlDS, macular degeneration, sorcening, diagnosis, therapeutch target identification and validation, and contain and contains an
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           genetic engineering, pharmacogenomics, studying gene function, and gene
mapping (e.g., of single nucleotide polymorphisms). The present sequence
represents the lower strand of a human TERC-targeted double-stranded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; sinA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; mixNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                        New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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100.0%; Pred. No. 1.5e+02;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 19 BP; 7 A; 4 C; 8 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human TERC siNA lower strand, SEQ ID 277.
                                                                                                                                                                                                                                                                                                                                Example 3; SEQ ID NO 275; 145pp; English.
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                                         2002US-0396782P.
2002US-0396600P.
2002US-0406784P.
                                                                                           2002US-0408378P
2002US-0409293P
                                                                                                                               15-JAN-2003; 2003US-0440129P
                          2002US-0363124P
                                                                                                                                                                     (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                      Beigelman L;
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Matches 19; Conservative
                                                                                                                                                                                                                                          WPI; 2003-689777/65.
                        11-MAR-2002;
06-JUN-2002;
17-JUL-2002;
29-AUG-2002;
05-SEP-2002;
09-SEP-2002;
                                                                                                                                                                                                          Mcswiggen J,
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonuclectide and an antisense oligonuclectide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonuclectides, and can be chemically synthesised, can contain deoxyribonuclectides, and can be chemically synthesised. Sequence of kits for the in vitro or in vivo delivery of siNA, conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue transplants for the treatment of a variety of conditions. They may be transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infections diseases (specifically cused for treating cancer, restenosis, infections diseases (specifically
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Pred. No. 1.5e+02;
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ID ADF93555 standard; RNA; 19 BP.
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05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409293P.
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2002US-0386782P.
2002US-0396600P.
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WO2003070742-A1.
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Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological;
                                        neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiathritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short halrpin RNA; shRNA; expression modulation; gene therapy; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
                                                                                                                                                                                                                                                                                                                                                                                            New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
          Human TERC siNA lower strand, SEQ ID 282
                                                                                                                                                                                                                                   2002US-0363124P.
2002US-0386782P.
2002US-0396600P.
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05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409293P.
15-JAN-2003; 2003US-0440129P.
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                                                                                                                                  Homo sapiens.
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06-JUN-2002;
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                                                                                                                                                                            28-AUG-2003.
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense credions, or alternatively are assembled from a sense onigonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short can contain deoxyribonucleotides, and can be chemically synthesised, can contain a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express sinha. The siNAs are used to modulate expression of the tellomerase genes in cells, tissue caphants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene companies the lower strand of a human TERC-targeted double-stranded Example 3; SEQ ID NO 282; 145pp; English.

Seguence 19 BP; 3 A; 8 C; 8 G; 0 T; 0 U; 0 Other;

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                      Gaps
                       ö
4.2%; Score 19; DB 1; Length 19; 00.0%; Pred. No. 1.5e+02;
                      0; Indels
       100.0%; Preq. ....
                       19; Conservative
Query Match
Best Local Similarity
                        Matches
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19 CTGCGGCGGGTCGCCTGCC 1

셤

RESULT 198

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Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antinfeumatory; agene therapy; telomerase; human; terc; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short interfering RNA; siRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
                                                      Human TERC siNA lower strand, SEQ ID 283
ADF93556 standard; RNA; 19 BP
                                                                                                                                                                                                                  11-FEB-2003; 2003WO-US004088
                                     26-FEB-2004 (first entry)
                                                                                                                                                                              WO2003070742-A1.
                                                                                                                                                            Homo sapiens.
                                                                                                                                                                                                                                     20-FEB-2002;
                                                                                                                                                                                                28-AUG-2003
                   ADF93556;
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(RIBO-) RIBOZYME PHARM INC. Mcswiggen J, Beigelman L;

2002US-0363124P. 2002US-0386782P. 2002US-0396600P.

11-MAR-2002; 06-JUN-2002;

17-JUL-2002; 29-AUG-2002;

2002US-0406784P. 2002US-0408378P. 2002US-0409293P.

05-SEP-2002; 15-JAN-2003;

2003US-0440129P

WPI; 2003-689777/65.

New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.

3xample 3; SEQ ID NO 283; 145pp; English.

The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense could be comprised and antisense could be stranded. They further comprise sense and antisense could be stranded. They further comprise sense and antisense contains and sold contains a sense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short contain deoxyribonucleotides, and can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised. Or relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are captains or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be transplants for the treatment of a variety of conditions. They may be conditions and rheumatoid arthritis. The siNAs are also useful for drug screening, disagnosis, luque srythematosus, AIDS, macular degeneration, genetic engineering, pharmacogenomics, studying gene function, and gene companying (e.g., of shalle mucleotide polymorphisms). The present sequence represents the lower strand of a human TERC-targeted double-stranded

Gaps

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ADF93291;

RESULT 199

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SINA.

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explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple solerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the upper strand of a human TERC-targeted double-stranded siNA, which is identical to the c-fos transcript target sequence.
                                                                                                                                                                                                                                                                                                                                                                                      Score 19; DB 1; Length 19;
Pred. No. 1.5e+02;
9; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                             Sequence 19 BP; 3 A; 4 C; 3 G; 0 T; 9 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               32 TGGCCATTTTTGTCTAAC 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 UGGCCAUUUUUUGUCUAAC 19
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                                                                                                                                                                                                                                                                                                                                                                                      ch 4.2%;
l Similarity 52.6%;
10; Conservative
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2002US-0386782P.
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2002US-0409293P.
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                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO2003070742-A1.
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06-JUN-2002;
17-JUL-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADF93302;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; tablomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; siRNA; couble-stranded RNA; micro-RNA; miRNA; short dargnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human TERC transcript target sequence/siNA upper strand, SEQ ID 8.
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                                                                                                      DB 1; Length 19;
1.5e+02;
                                                                                                                                                            0; Indels
                                             Sequence 19 BP; 0 A; 3 C; 13 G; 0 T; 3 U; 0 Other;
                                                                                                                    100.0%; Prea. ...
                                                                                                      Score 19;
Pred. No.
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                                                                                                                                                                                                         230 CCAGCCCCGAACCCGGC 248
                                                                                                                                                                                                                                                         19 CCAGCCCCGAACCCCGCC 1
                                                                                                                                                                                                                                                                                                                                                                                                       BP.
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2002US-0386782P.
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2002US-0408378P.
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                                                                                                      4.28;
                                                                                                                                                                                                                                                                                                                                                                                                 ADF93291 standard; RNA; 19
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                                                                                          Query Match
Best Local Similarity 100.0
Matches 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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17-JUL-2002;
29-AUG-2002;
05-SEP-2002;
09-SEP-2002;
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Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; sinA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shrap, expression medulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may
                                                                          Human TERC transcript target sequence/siNA upper strand, SEQ ID 19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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WPI; 2003-689777/65.

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be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the sings include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short harpin RNA (shRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically mythesised, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and vector or express sed. The invention also relates to kits for the invitro or in vivo delivery of siNA, conjugates and/or complexes of siNA, and vectors that express since siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and tended for treating cancer, restencesis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, ce.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, ergineering, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the upper strand of a human TERC-targeted double-stranded sink, which is identical to the c-fos transcript target sequence.
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100.0%; Pred. No. 1.5e+02;
tive 0; Mismatches 0; Indels
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2002US-0386782P.
2002US-0396600P.
2002US-0406784P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity 100.0
Matches 19; Conservative
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA convergulate expression of the one or more telomerase genes by RNA convergence. The siNAs may or may not comprise sense and antisense of interference. The siNAs may or may not comprise sense and antisense of ingoniclectide. Specifically, the siNAs inclored short interfering RNA (shRNA), double-stranded RNA, micro-RNA (miRNA) and short interfering RNA (shRNA). The siNAs can be unmodified or chemically synthesised, can contain deoxyribonuclectides, and can be chemically synthesised or chemically southesised conformally conjugates and/or complaxes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue captants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically strended diseases).

Control of a multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The sinAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, serion ceptoses the upper strand of a human TRRC-targeted double-stranded capture represents the upper strand of a human TRRC-targeted double-stranded cupier supper strand of a human TRRC-targeted double-stranded cupier sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gape
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human TERC transcript target sequence/siNA upper strand, SEQ ID 12.
                                                 New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 19 BP; 1 A; 2 C; 12 G; 0 T; 4 U; 0 Other;
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                                                                                                                        Example 3; SEQ ID NO 7; 145pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           14 TGGGCCTGGGAGGGGTGGT 32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 UGGGCCUGGGAGGGGUGGU 19
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11-MAR-2002; 2002US-0363124P.
06-JUN-2002; 2002US-0386782P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity 78.9
nes 15; Conservative
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Matches
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and antisense oligonucleotide. Specifically, the siNAs include abort interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA, conjugates and/or complexes of siNA, and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and cransplants for the treatment of a variety of conditions. They may be used for treating cancer restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, c.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the upper strand of a human TERC-targeted double-stranded sinAA, which is identical to the c-fos transcript target sequence.
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4.2%; Score 19; DB 1; Length 19;
Best Local Similarity 73.7%; Pred. No. 1.5e+02;
Matches 14; Conservative 5; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 19 BP; 2 A; 6 C; 6 G; 0 T; 5 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                     Example 3; SEQ ID NO 12; 145pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          104 CTCGCTGACTTTCAGCGGG 122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADF93545 standard; RNA; 19 BP
                         29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409293P.
2002US-0396600P
                                                                                                15-JAN-2003; 2003US-0440129P
                                                                                                                                                (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                Mcswiggen J, Beigelman L;
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siNAs), double-stranded RNA, micro-RNA (mirRNA) and short can contain deoxyribonucleotides, and can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, and/or complexes of siNA; and vectors that express in relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue caplants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treatsplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single mucleotide polymorphisms). The present sequence contains and part with a substance of the single mucleotide polymorphisms). The present sequence the single mucleotide polymorphisms). The present sequence the single mucleotide polymorphisms).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 3; SEQ ID NO 272; 145pp; English.
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                                                                                                                                                                              11-MAR-2002; 2002US-0363124F.
06-UTV-2002; 2002US-0386782F.
17-UTL-2002; 2002US-0396600F.
29-AUG-2002; 2002US-0406784F.
05-SEP-2002; 2002US-0408378F.
09-SEP-2002; 2002US-0408378F.
15-JAN-2003; 2003US-0440129F.
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ses 19; Conservative
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                      WO2003070742-A1.
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Matches
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ADF93561 standard; RNA; 19 BP

ADF93561/

ADF93561;

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The invention relates to short interfering nucleic acids (sinA) which downrequiate expression of the one or more telomerase genes by RNA interference. The sinAs may or may not comprise ribonucleotides and may be double or single stranded They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the sinAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA, and vectors that express sinA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and used for treating cancer, restencesis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, ADDS, macular degeneration, skin ulcers and rheumatoid arthitis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, agenetic engineering, pharmacogenomics, studying gene function, and sequence mapping (e.g., of single nucleotide polymorphisms). The present sequence mapping (e.g., of single nucleotide polymorphisms). The present sequence
Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; slNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
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2002US-0406784P.
2002US-0408378P.
2002US-0409293P.
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2002US-0386782P
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29-AUG-2002; 2
05-SEP-2002; 2
09-SEP-2002; 2
                                                                                                                                                                                                             Homo sapiens.
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06-JUN-2002;
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA conterference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs can be unmodified or charmacontain deoxyribonucleotides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA, conjugates and/or complexes of siNA, and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically problems).
                                                                                                                                                         Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short harpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 3; SEQ ID NO 288; 145pp; English.
                                                                                                                    Human TERC siNA lower strand, SEQ ID 288.
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2002US-0406784P.
2002US-0408378P.
2002US-0409293P.
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2002US-0386782P.
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                                                                             26-FEB-2004 (first entry)
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                                                                                                                                                                                                                                                                                                                                                        Homo sapiens.
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06-JUN-2002;
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Sequence 19 BP; 3 A; 9 C; 7 G; 0 T; 0 U; 0 Other;

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Gaps

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Length 19;

4.2%; Score 19; DB 1; Length 19; 100.0%; Pred. No. 1.5e+02; ive 0; Mismatches 0; Indels

266 CCGGGGCTTCTCCGGAGGC 284

8

Conservative

Local Similarity es 19; Conserv

Query Match Matches 19 cceegerrereceagee 1

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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonuclectide and an attisense oligonuclectide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonuclectides, and can be chemically synthesised. Expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue used to modulate expression of the telomerase genes in cells, tissue transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically
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                                                                                      Gaps
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                  Score 19; DB 1; Length 19;
Pred. No. 1.5e+02;
                                                                                  0; Indels
4.2%; Scc. 100.0%; Pred. No. ...
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                                                                                                                                                           320 CCGCGGGTCTCTCGGGGGC 338
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11-MAR-2002; 2002US-0363124P.

06-JUN-2002; 2002US-0386782P.

17-JUL-2002; 2002US-0396600P.

29-AUG-2002; 2002US-0406784P.

05-SEP-2002; 2002US-0408378P.

09-SEP-2002; 2002US-0408378P.
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                                                                                          Conservative
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                                                      Local Similarity
nes 19; Conserv
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                         Query Match
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protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, ALDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene rapping e.g., of single nucleoride polyancphisms). The present sequence represents the upper strand of a human TERC-targeted double-stranded siNA, which is identical to the c-fos transcript target sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflemmatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering nucleic acid; siNA; short interfering nucleic acid; siNA; micro-RNA; minNA; short interfering siRNA; attended RNA; micro-RNA; minNA; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
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                                                                                                                                                                                                           Score 19; DB 1; Length 19;
Pred. No. 1.5e+02;
5; Mismatches 0; Indels
                                                                                                                                                                           Sequence 19 BP; 2 A; 9 C; 3 G; 0 T; 5 U; 0 Other;
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                                                                                                                                                                                                                                                                                          140 GCCGCCTTCCACCGTTCAT 158
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2002US-0386782P.
2002US-0396600P.
2002US-0406784P.
2002US-0409293P.
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                                                                                                                                                                                                                                  Local Similarity 73.7
les 14; Conservative
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05-SEP-2002;
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interfering RNA (BiRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (BhRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in calls, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenoisis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lubus arythematosus AIDS, macular degeneration, eigen the understand rheumatoid arthritis. The siNAs are also useful for drug screnning, diagnosis, therapeutic target identification and validation, generated engapsing (e.g., of single nucleotide polymorphisms). The present sequence represents the upper strand of a human TERC-targeted double-stranded
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 356 CTTTCAGGCCGCAGGAGA 374
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2002US-0386782P.
2002US-0396600P.
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2002US-0408378P
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15-JAN-2003; 2003US-0440129P
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29-AUG-2002;
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06-JUN-2002;
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New short interfering nucleic acid downregulates expression of the

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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA comprise reference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense of oligonucleotide. Specifically, the siNAs include and antisense contain deoxyribonucleotides, and can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be enemically synthesised. Components of solver the invitro or in vivo delivery of siNA; conjugates can contain deoxyribonucleotides, and can be enemically synthesised. The invention also can contain deoxyribonucleotides, and can be chemically synthesised. Complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue cused to modulate expression of the telomerase genes in cells, tissue cused to modulate expression of the telomerase genes in cells, tissue cused to modulate expression of the vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treatment calcetion, or autoimmune or age-related diseases. Confections diseases (specifically processed in ulciple sclerosis, lupue erythematosus, AIDS, macular degeneration, sind genes compliants diagnosis, therapeutic rarget famentian and validation. Skin ulcers and rheumatoid arthritis. The sinNas are also useful for drug capteric engineering, pharmacogenomics, studying gene function, and gene captessents the upper strand of a human TERC-targeted double-stranded cusples sclerosis.
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telomerase gene useful e.g. for treatment and diagnosis of cancer.
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                                         Example 3; SEQ ID NO 27; 145pp; English.
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2002US-0386782P.
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2002US-0408378P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity 94.7%;
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADF93292 standard; RNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  26-FEB-2004 (first entry)
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29-AUG-2002;
05-SEP-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       28-AUG-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 209
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA committees and antisense interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short interfering RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue caplants or organisms (e.g., by ex vivo gene therapy), or in grafts and treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically proteozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lugus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug correpting, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene corpusation the upper strand of a human TERC-targeted double-stranded creptered to the c-fos transcript target sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering uncleic acid; siNA; short interfering RNA; siRRA; double-stranded RNA; micro-RNA; miRNA; short interfering RNA; siRRA; double-stranded RNA; micro-RNA; miRNA; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human TERC transcript target sequence/siNA upper strand, SEQ ID 15.
                                                                                                                                                             telomerase gene useful e.g. for treatment and diagnosis of cancer.
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0
                                                                                                                                            New short interfering nucleic acid downregulates expression of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             4.2%; Score 19; DB 1; Length 19;
84.2%; Pred. No. 1.5e+02;
tive 3; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 19 BP; 5 A; 5 C; 6 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                           Example 3; SEQ ID NO 9; 145pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               50 CCCTAACTGAGAAGGGCGT 68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 CCCUAACUGAGAAGGGCGU 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADF93298 standard; RNA; 19 BP
 09-SEP-2002; 2002US-0409293P.
                15-JAN-2003; 2003US-0440129P.
                                                (RIBO-) RIBOZYME PHARM INC.
                                                                              Mcswiggen J, Beigelman L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             26-FEB-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 84.2%
                                                                                                             WPI; 2003-689777/65.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADF93298;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 210
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the inversious trainers to smort interioring nucercy dense by RNA interference. The sinks may or may not comprise sense and antisense conductor. The single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and antisense conductoring the Specifically, the sinks include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short can contain deoxyribonucleotides, and can be chemically synthesised, can complexes of sink, and vector or invivo delivery of sink, conjugates and/or complexes of sink, by ex vivo genes in cells, tissue cused to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be corpused for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, ce.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The sinks are also useful for drug screening, diagnosis, therapeutic target identification and validation, capanesing (e.g., of single mucleotide polymorphisms). The present sequence capanes, which is identical to the c-foe transcript target sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to short interfering nucleic acids (siNA) which
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4.2%; Score 19; DB 1; Length 19;
Best Local Similarity 84.2%; Pred. No. 1.5e+02;
Matches 16; Conservative 3; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 19 BP; 11 A; 3 C; 2 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 3; SEQ ID NO 15; 145pp; English.
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                                                                                                                                                                                                                                                                                                                      17-JUL-2002; 2002US-0396600P.
29-AUG-2002; 2002US-04067B4P.
05-SEP-2002; 2002US-040837BP.
09-SEP-2002; 2002US-040837BP.
                                                                                                                                                                                                                             2002US-0363124P.
2002US-0386782P.
                                                                                        11-FEB-2003; 2003WO-US004088.
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                                                                                                                                                                                      20-FEB-2002;
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28-AUG-2003
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US09436060A-16.rng.sl

Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; mioro-RNA; miRNA; short interfering RNA; shrNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.

Human TERC transcript target sequence/siNA upper strand, SEQ ID 30

26-FEB-2004 (first entry)

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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense of ligonucleotide. Specifically, the siNAs include short interfering RNA (shRNA), double-stranded RNA, micro-RNA (miRNA) and short interfering RNA (shRNA). The siNAs can be unmodified or chemically synthesised. Can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised. Cexpressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue cexplants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically protected in ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeuth crayer identification and validation, shin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeuth crayer identification and validation, sereening, diagnosis, therapeuth expecting servents the upper strand of a human TERC-targeted double-stranded crepresents the upper strand of a transcript target sequence.
RNA interference; short interfering nucleic acid; siNA; short interfering RNA; double-stranded RNA; micro-RNA; miRNA; short interfering RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 3; SEQ ID NO 28; 145pp; English.
                                                                                                                                                                                                                                                                                                                                             2002US-0386782P.
2002US-0396600P.
2002US-0406784P.
2002US-0408378P.
2002US-0409293P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mcswiggen J, Beigelman L;
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                                                                                                                                                                          WO2003070742-A1.
                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                              11-FEB-2003;
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05-SEP-2002;
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                                                                                                                                                                                                                                                                                                                                                    06-JUN-2002;
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2002US-0396600P. 2002US-0406784P. 2002US-0408378P. 2002US-0409293P.

05-SEP-2002; 15-JAN-2003;

2003US-0440129P

(RIBO-) RIBOZYME PHARM INC.

Mcswiggen J, Beigelman L; WPI; 2003-689777/65.

2002US-0363124P

11-FEB-2003; 2003WO-US004088

20-FEB-2002; 06-JUN-2002; 17-JUL-2002; 29-AUG-2002;

11-MAR-2002

WO2003070742-A1. Homo sapiens

28-AUG-2003

New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.

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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise stonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and an interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, can complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue used to complexe of siNA; and vectors that express siNA. The siNAs are used to complexe of siNA; and vectors that express siNA. The siNAs are used to complexe of siNA; and vectors that express siNA. The siNAs are used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus expthematosus, ANDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug creening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence is read of a human TERC-targeted double-stranded contains and the upper strand of a human TERC-targeted double-stranded contains the presents that the upper strand of a human TERC-targeted double-stranded contains and the upper strand of a human TERC-targeted double-stranded double-stranded degeneration and sequence in the upper strand of a human transfer and restranded deserves the superior of the tearther of the superior of the stranded deserves the superior of the superi
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Pred. No. 1.5e+02;
2; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             392 CGCGCGCGCGCGATTCCC 410
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Best Local
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siNA, which is identical to the c-fos transcript target sequence.
                                                                                   Length 19;
                                      Sequence 19 BP; 5 A; 8 C; 4 G; 0 T; 2 U; 0 Other;
                                                                                   Score 19; DB 1; 1
Pred. No. 1.5e+02;
                                                                                   4.2%;
                                                                                   Query Match
Best Local Similarity
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ADF93313 standard; RNA; 19 BP.

RESULT 212

충 셤 ADF93313 ID ADF9

US09436060A-16.rng.sl

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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide an antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised.

Expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue used to modulate expression of the telomerase genes in cells, tissue transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically corrocal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The sinAs are also useful for drug
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  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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0
0; Indels
2; Mismatches
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                                                                                                                                                                                                                                                                                                                          Human TERC siNA lower strand, SEQ ID 284.
                                        428 ACCCAGGACTCGGCTCACA 446
                                                                                  1 Acceadadoucedeucaca 19
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2002US-0406784P.
2002US-0408378P.
2002US-0409293P.
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2002US-0363124P.
2002US-0386782P.
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                                                                                                                                                                                           ADF93557 standard; RNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (RIBO-) RIBOZYME PHARM INC.
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                                                                                                                                                                                                                                                                               (first entry)
  17; Conservative
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06-JUN-2002;
17-JUL-2002;
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05-SEP-2002;
09-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
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                                                                                                                                                                                                                                                                               26-FEB-2004
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                                                                                                                                                                                                                                     ADF93557;
                                                                                                                                                   RESULT 213
Matches
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                          genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the lower strand of a human TERC-targeted double-stranded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               vasotropic; protozoacide; immunosuppressive; dermatological;
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screening, diagnosis, therapeutic target identification and validation,
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100.0%; Pred. No. ...
0; Mismatches
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2002US-0363124P.
2002US-0386782P.
2002US-0406784P.
2002US-040878P.
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                                                                                                                                                                                                                                                          Query Match 4.2
Best Local Similarity 100.
Matches 19; Conservative
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06-JUN-2002;
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expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of sinh, conjugates and/or complexes of sinh, and vectors that express sinh. The sinhs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenced, infectious diseases (specifically protozcal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNhs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nuclectide polymorphisms). The present sequence represents the lower strand of a human TERC-targeted double-stranded
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Pred. No. 1.5e+02;
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                                                                                                                                                                                                                                                                                            Sequence 19 BP; 2 A; 7 C; 9 G; 0 T; 1 U; 0 Other;
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100.0%; Pred. No. ...
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29-AUG-2002; 2002US-0406784P-
0S-SEP-2002; 2002US-0408378P-
0S-SEP-2002; 2002US-0409293P-
1S-JAN-2003; 2003US-0440129P-
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2002US-0386782P.
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19; Conservative
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Best Local S
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Example 3; SEQ ID NO 278; 145pp; English.

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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and an interfering RNA (siNAA), double-stranded RNA, micro-RNA (miRNA) and short niterfering RNA (siNAA). Adouble-stranded RNA, micro-RNA (miRNA) and short niterfering RNA (siNAA). The siNAS can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically sorthesised, cancomplexes of siNA, and vectors that express siNA. The siNAS are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be
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                                                                                                                                                                                                                                                                                                                             used for treating cancer, restenosis, infectious diseases (apportifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the lower strand of a human TERC-targeted double-stranded
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100.0%; Pred. No. 1.5e+02;
ive 0; Mismatches 0;
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2002US-0363124P.
2002US-0386782P.
2002US-0396600P.
2002US-0406784P.
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2002US-0409293P.
2003US-0440129P.
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les 19; Conserv
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11-MAR-2002;
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2002US-0363124P. 2002US-0386782P. 2002US-0396600P. 2002US-0406784P. 2002US-0408378P.

09-SEP-2002; 2002US-0409293P. 15-JAN-2003; 2003US-0440129P. (RIBO-) RIBOZYME PHARM INC. Mcswiggen J, Beigelman L; New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.

WPI; 2003-689777/65.

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20-FEB-2002;
11-MAR-2002;
06-JUN-2002;
                                                                    17-JUL-2002;
29-AUG-2002;
05-SEP-2002;
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                                                                                                        New short interfering nucleic acid downregulates expression of the
                                                                                                                          telomerase gene useful e.g. for treatment and diagnosis of cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 19 BP; 1 A; 5 C; 6 G; 0 T; 7 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     7; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human TERC siNA lower strand, SEQ ID 273.
                                                                                                                                                            Example 3; SEQ ID NO 10; 145pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 UAGGCGCCGUGCUUUGCU 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        68 TAGGCGCCGTGCTTTTGCT 86
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADF93546 standard; RNA; 19 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    63.2%;
(RIBO-) RIBOZYME PHARM INC.
                                   Mcswiggen J, Beigelman L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      26-FEB-2004 (first entry)
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Matches 12; Conservative
                                                                    WPI; 2003-689777/65.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 217
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADF93546/
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The inversion transfer to such the one or more telomerase genes by NRA interference. The siNAs may or may not comprise sense and antisense be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and antisense regions, or alternatively are assembled from a sense oligonucleotide and antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siNRA), double-stranded RNA, micro-RNA (miRNA) and short can contain deoxyribonucleotides, and can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised. Correlates to kits for the in vitro or in vivo delivery of siNR, conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue caplants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be concerned to the treatment of a variety of conditions. They may be corrected in the treatment of a variety of conditions. They may be corrected or multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, disagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene constraints the lower strand of a human TERC-targeted double-stranded
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                                                                        The invention relates to short interfering nucleic acids (siNA) which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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100.0%; Pred. No. 1.5e+02;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 19 BP; 3 A; 6 C; 5 G; 0 T; 5 U; 0 Other;
Example 3; SEQ ID NO 273; 145pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19 cccraacreagaagacer 1
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ADF93562/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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11-FEB-2003; 2003WO-US004088.

Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; meuroprotective; anti-HIV, ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shrNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.

Human TERC transcript target sequence/siNA upper strand, SEQ ID 25

(first entry)

26-FEB-2004

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The invention relates to short interfering nucleic acids (siNA) which downrequiate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short interfering RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA, conjugates and/or complexes of siNA, and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in calls, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and trensplants for the treatment of a variety of conditions. They may be used for treating cancer, restenoisis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lubus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthitis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, agenetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence
drug screening; diagnosis; therapeutic target identification;
pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
                                                                                                                                                                                                                                                                                                                                                            New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4.2%; Score 19; DB 1; Length 19; 100.0%; Pred. No. 1.58+02; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 19 BP; 2 A; 9 C; 5 G; 0 T; 3 U; 0 Other;
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                                                                                                                                                                           06-JUN-2002; 2002US-0386782P
17-JUL-2002; 2002US-039660P
29-AUG-2002; 2002US-0406784P
05-SEP-2002; 2002US-0409378P
09-SEP-2002; 2002US-0409293P
                                                                                                                       11-FEB-2003; 2003WO-US004088
                                                                                                                                                                2002US-0363124P
                                                                                                                                                                                                                                                  15-JAN-2003; 2003US-0440129P
                                                                                                                                                                                                                                                                              (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                         Beigelman L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                     WPI; 2003-689777/65.
                                                                  WO2003070742-A1
                                         Homo sapiens.
                                                                                                                                                                                                                                                                                                         Mcswiggen J,
                                                                                                                                                                 11-MAR-2002;
                                                                                             28-AUG-2003
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2002US-0386782P. 2002US-0396600P. 2002US-0406784P. 2002US-0408378P. 2002US-0409233P.

29-AUG-2002; 05-SEP-2002; 09-SEP-2002; 15-JAN-2003; (RIBO-) RIBOZYME PHARM INC.

Mcswiggen J, Beigelman L;

WPI; 2003-689777/65.

2002US-0363124P

20-FEB-2002;

11-MAR-2002; 06-JUN-2002;

11-FEB-2003; 2003WO-US004088

WO2003070742-A1. Homo sapiens.

28-AUG-2003

New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.

Example 3; SEQ ID NO 25; 145pp; English.

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The invention relates to short interfering muclaicate actus (shink) which convergulate expression of the one or more telomerase genes by RNA interference. The sinks may or may not comprise ribonucleotides and maisense be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the sinks include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (mRNA) and short hairpin RNA (siRNA). The sinks can be unmodified or chemically withheaised, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA, conjugates and/or complexes of siNA, and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue captures or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplant rejection, or autoimmune or age-related diseases, protozoal), transplant rejection, or autoimmune or age-related diseases, protozoal), transplant rejection, or autoimmune or age-related diseases, comultiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumacoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, barmacogenomics, studying gene function, and genetic engenetation and genetic engenetation and genetic engenetation in the upper strand of a human TERC-targeted double-stranded diseases.
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invention relates to short interfering nucleic acids (siNA) which
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  siNA, which is identical to the c-fos transcript target sequence.
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Best Local Similarity 89.59
Watches 17, Conservative
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Gaps

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338 CGAGGCCGAGGTTCAGGCC 356

Conservative

19;

Matches

CGAGGGGAGGTTCAGGCC 1

13

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ADF93308 standard; RNA; 19 BP

RESULT 219

ADF93308

ADF93308
ID ADF9
XX
AC ADF9
XX

338 CGAGGCGAGGTTCAGGCC 356

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represents the lower strand of a human TERC-targeted double-stranded

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Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
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                                                                                                                                                                                                                                                                                                                                                        Human TERC siNA lower strand, SEQ ID 279.
BP.
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2002US-0386782P.
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29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
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                                                                                                                                                                             ADF93552 standard; RNA; 19
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                                                                                                                                                                                                                                                                                                   (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2003-689777/65.
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06-JUN-2002;
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                                                                                                                                                                                                                                        ADF93552;
                                                                                                                        RESULT 220
                                                                                                                                              ADP 93552/
XX ADP 9
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonuclectide and antisense oligonuclectide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short interfering RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonuclectides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates confaint expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and candor complants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rhemamatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, centured mapping (e.g., of single nucleotide polymorphisms). The present sequence
Example 3; SEQ ID NO 279; 145pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                        Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; sink; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; mikna; short hairpin RNA; shrNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs can be unmodified or chemically modified, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are
                                                                                                                       Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 telomerase gene useful e.g. for treatment and diagnosis of cancer.
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                                                                                    Length 19;
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                                                Sequence 19 BP; 3 A; 2 C; 3 G; 0 T; 11 U; 0 Other;
                                                                                   DB 1; Le
1.5e+02;
                                                                  4.2%; Scor.
100.0%; Pred. No. 1...
... 0; Mismatches
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                                                                                                                                                    158 TTCTAGAGCAAACAAAAA 176
                                                                                                                                                                                    Trcragagcaaacaaaaa 1
                                                                                                                                                                                                                                                                         BP.
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2002US-0386782P.
2002US-0396600P.
2002US-0406784P.
2002US-0408378P.
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                                                                                                                                                                                                                                                                         ADF93566 standard; RNA; 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mcswiggen J, Beigelman L;
                                                                           Query Match
Best Local Similarity 100.
Matches 19; Conservative
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06-JUN-2002;
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05-SEP-2002;
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used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenced in infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siMAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the lower strand of a human TERC-targeted double-stranded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; abhort hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERT; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA
                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human TERC transcript target sequence/siNA upper strand, SEQ ID 13.
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                                                                                                                                                                                                                                                                                             Score 19; DB 1; Length 19;
Pred. No. 1.5e+02;
                                                                                                                                                                                                                                                                                                                                   0; Indels
                                                                                                                                                                                                                                                       Sequence 19 BP; 4 A; 8 C; 4 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                           Query Match 4.2%; Score 19; DB Best Local Similarity 100.0%; Pred. No. 1.5 Matches 19; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                          410 CTGAGCTGTGGGACGTGCA 428
                                                                                                                                                                                                                                                                                                                                                                                                  19 CTGAGCTGTGGGACGTGCA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADF93296 standard; RNA; 19 BP
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05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409293P.
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2002US-0386782P.
2002US-0396600P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mcswiggen J, Beigelman L;
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06-JUN-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 222
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interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short an antisense oligonucleotide. Specifically, the siNAs include short hard siRNA), double-stranded RNA, micro-RNA (miRNA) and short can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised. Confirm a vector or enzymatically synthesised. The invention also capteses of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue cand/or complexes of siNA; and vectors that express siNA. The siNAs are used to reganisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically proposed for conditions and rheumatoid arthritis. The siNAs are also useful for drug skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug cancer, cancer and signification and sense since the contains, diagnosis, therappeutic farget identification and validation, containing, diagnosis, therappeut captend for sense function and sense function and sense function.
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                                                                                                                                                                                                                                                                                                                                                                                                                    genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the upper strand of a human Tebreserts double-stranded siNs, which is identical to the c-fos transcript target sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4.2%; Score 19; DB 1; 189.5%; Pred. No. 1.5e+02; tive 2; Mismatches 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            122 GCGGAAAAGCCTCGGCCTG 140
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11-MMX-2002; 2002US-0363124P.
06-UIN-2002; 2002US-0386782P.
17-UUL-2002; 2002US-0396600P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0409373P.
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Best Local Similarity 89.5
Matches 17; Conservative
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                                                                                                                          The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA characterence. The siNAs may or may not comprise ribonuclecides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs can be unmodified or chemically modified, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA, and vectors that express sinc. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restencisis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus. AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, agenematic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the lower strand of a human TERC-targeted double-stranded
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                                                New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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100.0%; Pred. No. 1.5e+02;
ive 0; Mismatches 0; Indels
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                                                                                                   Example 3; SEQ ID NO 271; 145pp; English.
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11-MAR-2002; 2002US-0363124P.
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Best Local Similarity 100.(
Matches 19; Conservative
                 WPI; 2003-689777/65
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNas may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and antisense oligonucleotide. Specifically, the siNas include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short interfering RNA (shRNA). The siNas can be unmodified or chemically modified, can contain decoxribbonucleotides, and can be chemically synthesised, can contain decoxribbonucleotides, and can be chemically synthesised, can contain decoxribbonucleotides, and can be chemically synthesised, can contain decoxribbonucleotides, and vectors that express sink. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are cused to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be cotozeal), transplant rejection, or autoimmune or age-related diseases, cells multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening diagnosis, therapeutic target identification and validation, can mapping (e.g., of single mucleotide polymorphisms). The present sequence mapping (e.g., of single mucleotide polymorphisms). The present sequence constants the lower strand of a human TERC-targeted double-stranded
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100.0%; Pred. No. 1.5e+02;
tive 0; Mismatches 0; Indels
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                                   17-JUL-2002; 2002US-0396600P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409293P.
06-JUN-2002; 2002US-0386782P.
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Human; ss; SDSO; short double stranded oligonucleotide; cleavage site; viral infection; malignant tumour; genetic disease; metabolic disease; gene chip; protein chip; microarray; gene drug; Dermogene; Lungene; Hepatogene; Leukogene; Lymbhogene; Prostogene; Braatogene; Braatogene; Braintumogene; Skin-whitogene; short interfering RNA; sIRNA; cancer;

RNA interference

US2004072769-A1

15-APR-2004

Homo sapiens

16-SEP-2002; 2002US-00016490. 16-SEP-2002; 2002US-00016490.

WPI; 2004-355427/33

(YINJ/) YIN J Q.

Yin JQ;

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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonuclectide and an interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonuclectides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in calls, tissue explants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically cortozoal), transplant rejection, or autoimmune or age-related diseases, es multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the lower strand of a human TBRC-targeted double-stranded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gapa
                                                                                                                                                                                                                                                                          New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 19; DB 1; Length 19;
Pred. No. 1.5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 19 BP; 1 A; 3 C; 10 G; 0 T; 5 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%; Pred. ...
                                                                                                                                                                                                                                                                                                               Example 3; SEQ ID NO 286; 145pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 284 CACCCACTGCCACCGCGAA 302
                                                                                                                     2002US-0386782P
                                                                                                                              2002US-0396600P
2002US-0406784P
                                                                                                                                                      2002US-0408378P
                                                                      11-FEB-2003; 2003WO-US004088
                                                                                                       2002US-0363124P
                                                                                                                                                                              2003US-0440129P
                                                                                             2002US-0358580P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4.28;
                                                                                                                                                                                                     (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                             Mcswiggen J, Beigelman L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 100.
Matches 19; Conservative
                                                                                                                                                                                                                                                     WPI; 2003-689777/65.
                      WO2003070742-A1.
                                                                                                                                                      05-SEP-2002; 2
09-SEP-2002; 2
15-JAN-2003; 2
 Homo sapiens.
                                                                                                                    06-JUN-2002;
                                                                                                                              17-JUL-2002;
                                                                                            20-FEB-2002;
                                                                                                      11-MAR-2002;
                                                                                                                                            29-AUG-2002;
                                              28-AUG-2003
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Sucrement. Also included are plantmaceutical compositions of gene drugs councilled. As becaused are plantmaceutical compositions of gene drugs of such as Dermogene, Lungene, Hepatogene, Leukogene, Lymphogene, Prostogene, Breastogene, Braintumogene and Skin-whitogene including but composed to the part or all of the following components: single or a group of specific 19-25 at daRNA, 19-25 at daRNA and/or bingle-stranded RNA and/or DNA with the special pattern, 5'-CGGAT(U)-3' or its derivatives, one or more nucleic acid condensation agents (or none), one or more pharmaceutical carriers, one or more pattern active agents and additional specific cell-targeting proteins and other active agents and additional materials) and a simplified method for predicting and selecting as specific and efficacious small double-stranded oligonucleotides (SDSO), antisense oligonucleotide molecules or short interfering RNA (sIRNA) (comprising identifying a special pattern that can be localised in any position of an oligonucleotide sequence valuating the specificity of a selected sequence). The Short interfering RNA (sIRNA) are targeted against genes involved in viral infection, malignant tumours, genetic and short double-stranded oligonucleotides as a gene drug that can specifically inactivate a group of corresponding genes. The composition can's be used for treating diseases or disorders associated with abnormal expression of genes in cells or tissues of humans or animals, such as viral infections, cancer, or genetic or metabolic diseases. The present conversed or sequence is a target region for an SDSO from an human cDNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to screening, identifying or predicting, and assembling 19-25 nt double-stranded oligonucleotides (termed short double stranded oligonucleotides, SDSO) as active pharmaceutical compositions for the treatment of viral infections, malignant tumours, and genetic and metabolic diseases, comprising screening and identifying a specific and protein chip microarrays. The above method comprises screening the cisease-causing genes, over-expressing in cells and/or tissues, with the disease-causing genes, over-expressing in cells and/or tissues, with the gene chip and protein chip microarrays, identifying a specific DNA generic within the abnormal gene encoding a protein or playing other biological roles with the assistance of computer and specific software, predicting efficacious 19-25 nt double-stranded oligonucleotides with a 5'-AU(T)CCG-3' or 5'-U(T)CCG-3' special pattern complementary to at least a portion of an RNA molecule and making sure that selected sequence is not localised within the stem-loop of target mRNA with any related control of the pharmaceutical compositions of gene drugs of the pharmaceutical compositions of gene drugs
Designing and selecting short double-stranded oligonucleotides for treating viral infections, cancer and genetic or metabolic diseases, comprises using gene chip and protein chip microarrays to identify
                                                                                                                                                                                                                                                                                                                                                     Claim 12; SEQ ID NO 8; 58pp; English
                                                                                                                                                                                                       specific DNA sequences.
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Sequence 19 BP; 6 A; 4 C; 8 G; 1 T; 0 U; 0 Other;

4.2%; Score 19; DB 1; Length 19;

Query Match

Human telomerase RNA gene, SDSO target region #4.

(first entry)

01-JUL-2004

ADO22919;

BXXXXEXB

19 cacccacreccaccecaa 1

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ADO22919 standard; cDNA; 19 BP

RESULT 226 AD022919 antisense Oligonucleotide molecules or short interfering RNA (siRNA) (comprising identifying a special pattern that can be localised in any position of an oligonucleotide sequence evaluating the specificity of a selected sequence). The Short interfering RNA (siRNA) are targeted against genes involved in viral infection, malignant tumours, genetic an metabolic diseases. The methods are useful for designing and selecting short double-stranded oligonucleotides as a gene drug that can specifically inactivate a group of corresponding genes. The composition may be used for treating diseases or disorders associated with abnormal expression of genes in cells or tissues of humans or animals, such as viral infections, cancer, or genetic or metabolic diseases. The present sequence is a target region for an SDSO from an human cDNA.

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specific and efficacious small double-stranded oligonucleotides (SDSO),

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The invention relates to screening, identifying or predicting, and assembling 19-25 nt double-stranded oligonucleotides (termed short double stranded oligonucleotides, SDSO) as active pharmaceutical compositions for the treatment of viral infections, malignant tumours, and genetic and metabolic diseases, comprising screening and identifying a specific DNA sequence in an abnormal gene encoding a protein with gene chip and protein chip microarrays. The above method comprises screening the disease-causing genes, over-expressing in cells and/or tissues, with the abnormal gene encoding a protein or playing other compression of protein chip microarrays, identifying a specific DNA composition of protein chip microarrays, identifying a specific DNA composition of predicting efficacious 19-25 nt double-stranded oligonucleotides with a composition of predicting efficacious 19-25 nt double-stranded oligonucleotides with a composition of an RNA molecule and making sure that selected sequence is not localised within the stem-loop of target mRNA with any related software. Also included are pharmaceutical compositions of gene drugs (such as Dermogene, Dungene, Hepatogene, Leukogene, Lymphogene, Composition of specific 19-25 nt daRNA, 19-25 nt sRNA-cDNA, 19-25 nt daRNA and/or single-stranded RNA and/or NNA with the special pattern, 5-composition of specific oper more pharmaceutical carriers, one or more composition and additional agents (or none), one or more pharmaceutical carriers, one or more specific cell-targeting proteins and other active agents and additional materials) and a simplified method for predicting and selecting a
                      ö
                                                                                                                                                                                                                                                                                                                                                                                  Human; ss; SDSO; short double stranded oligonucleotide; cleavage site; viral infection; malignant tumour; genetic disease; metabolic disease; gene chip; protein chip; microarray; gene drug; Dernogene; Lungene; Hepatogene; Lungene; Lungene; Lungene; Lungene; Lungene; brostogene; Breastogene; Braintumogene; Skin-whitogene; short interfering RNA; siRNA; cancer;
                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Designing and selecting short double-stranded oligonucleotides for treating viral infections, cancer and genetic or metabolic diseases, comprises using gene chip and protein chip microarrays to identify
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                    Indels
                                                                                                                                                                                                                                                                                                                                            Human telomerase RNA gene, SDSO target region #3.
No. 1.5e+02;
natches 0;
                    0; Mismatches
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                                                                 372 AGAGGAACGGAGCGAGTCC 390
                                                                                                         1 AGAGGAACGGAGCGAGTCC 19
                                                                                                                                                                                                                ADO23063 standard; cDNA; 19 BP.
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100.08;
                                                                                                                                                                                                                                                                                                      (first entry)
  Best Local Similarity 100.
Matches 19; Conservative
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                                                                                                                                                                                                                                                             ADO23063;
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                                                                                                                                                                        RESULT 227
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Gaps

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0; Indels

Score 19; DB 1; Length 19; Pred. No. 1.5e+02;

Query Match 4.2%; Score 19; DB Best Local Similarity 100.0%; Pred. No. 1.5 Matches 19; Conservative 0; Mismatches

424 GTGCACCCAGGACTCGGCT 442 |||||||||||||||| 1 GTGCACCCAGGACTCGGCT 19

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Sequence 19 BP; 3 A; 7 C; 6 G; 3 T; 0 U; 0 Other;

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The invention relates to screening, identifying or predicting, and assembling 19-25 nt double-stranded oligonucleotides (termed short double stranded oligonucleotides, SDSO) as active pharmaceutical compositions for the treatment of viral infections, malignant tumours, and genetic and metabolic diseases, comprising screening and identifying a specific BNA sequence in an abnormal gene encoding a protein with gene chip and protein chip microarrays. The above method comprises screening the disease-causing genes, over-expressing in cells and/or tissues, with the
                                                                                                                                                                          Human; ss; SDSO; short double stranded oligonucleotide; cleavage site; viral infection; malignant tumour; genetic disease; metabolic disease; gene chip; protein chip; microarray; gene drug; Dernogene; Lungene; Hepatogene; Lunkogene; Lymbhogene; Prostogene; Breastogene; Braintumogene; Skin-whitogene; short interfering RNA; siRNA; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Designing and selecting short double-stranded oligonucleotides for treating viral infections, cancer and genetic or metabolic diseases, comprises using gene chip and protein chip microarrays to identify
                                                                                                                                           Human telomerase RNA gene, SDSO target region #5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 1; Page 19; 58pp; English.
                                  ADO23064 standard; cDNA; 19 BP
                                                                                                                                                                                                                                                                                                                                                                                                                         16-SEP-2002; 2002US-00016490.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              16-SEP-2002; 2002US-00016490.
                                                                                                            01-JUL-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      specific DNA sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2004-355427/33.
                                                                                                                                                                                                                                                                           RNA interference
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                                                                                                                                                                                                                                                                                                                  Homo sapiens.
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                                                                       ADO23064;
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RESULT 228
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WPI; 2004-355427/33.

gene chip and protein chip microarrays, identifying a specific DNA sequence within the abnormal gene encoding a protein or playing other biological rools with the assistance of computer and specific software, predicting efficacious 19-25 nt double-stranded oligonuclectides with a sequence is '-AU(T)CCG-3' or 5'-U(T)CCG-3' special pattern complementary to at sequence is not localised within the stem-loop of target mRNA with any related software. Also included are pharmaceutical compositions of gene drugs (such as Dermogene, Lungene, Leukogene, Lunghogene, Europane, Lungene, Europane, Lunghogene, Dermogene, Lungene, Dermogene, Lunghogene, Dermogene, Lunghogene, Dermogene, Lunghogene, Dermogene, Lunghogene, Dermogene, Lunghogene, Dermogene, Dermogene, Lunghogene, Dermogene, Lunghogene, Dermogene, Lunghogene, Dermogene, Lunghogene, Dermogene, Lunghogene, Dermogene, Dermo

Sequence 19 BP; 1 A; 2 C; 12 G; 4 T; 0 U; 0 Other;

Score 19; DB 1; Length 19; Pred. No. 1.5e+02; 0; Mismatches 0; Indels 4.2%; Sc._ 100.0%; Pred 0; M 1 TGGGCCTGGGAGGGGTGGT 19 14 TGGGCCTGGGAGGGGTGGT 32 19; Conservative Query Match Best Local Similarity Matches 용 ò

0; Gaps

ADO23062 standard; cDNA; 19 BP AD023062; RESULT 229 AD023062 XXX AC AD02 XXX XXX XXX XXX Hume XXX Hume XXX Hope Hope Hope XXX Hope Hope Home XXX Hope Home XXX Home Hope Home Hope XXX Home Hope Home Hope Home Home Hope Hope

01-JUL-2004 (first entry)

Human telomerase RNA gene, SDSO target region #2.

Human; 88; SDSO; short double stranded oligonucleotide; cleavage site; viral infection; malignant tumour; genetic disease; metabolic disease; gene chip; protein chip; microarray; gene drug; Dermogene; Lungene; Hepatogene; Lunkogene; Lymbhogene; Prostogene; Braintumogene; Skin-whitogene; short interfering RNA; siRNA; cancer; RNA interference

Homo sapiens.

US2004072769-A1

15-APR-2004.

16-SEP-2002; 2002US-00016490.

16-SEP-2002; 2002US-00016490.

(YINJ/) YIN J Q.

Yin JO;

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The invention relates to screening, identifying or predicting, and casembling 19-25 in double-stranded oligomucleotides (termed short double stranded oligomucleotides) assembling 19-25 in double-stranded oligomucleotides (termed short double for the treatment of viral infections, malignant tumours, and genetic and for the treatment of viral infections, malignant tumours, and genetic and control and apportant gene encoding a protein with gene chip and protein chip microarrays. The above method comprises screening the protein chip microarrays. The above method comprises screening the protein chip microarrays. The above method comprises screening the protein chip microarrays. The above method comprises screening the protein chip microarrays in cells and/or tissues, with the case and protein chip microarrays, identifying a specific DNA sequence within the abormal gene encoding a protein or protein chip microarrays. Include a specific base causing genes over-expession in cells and/or there is continued to a nRAA molecule and making sure that selected sequence of predicting efficacious 19-25 nt double-stranded oligonucleotides with a signal and protein of an RNA molecule and making sure that selected sequence of some to localised within the stem-loop of target manks with any related conference. Indeed are pharmaceutical compositions of gene drugs software. Also included are pharmaceutical compositions of gene limited to part or all of the following components: single or protein and or indeed the mand/or DNA with the special pattern, 5.

Cromyrials) and efficacious small double-stranded oligonucleotides (SDSO), and estimate and elimited to parameter manded composition of an oligonucleotide molecules or short interfering RNA (siRNA) are cargeted sequence. The methods are useful for designing and selecting manded oligonucleotide molecules or short interfering the segment of an oligonucleotide sequence evaluating the specific and efficacious small device as a gene drug that can be composition of an oligonucleotide molecu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                            Designing and selecting short double-stranded oligonucleotides for treating viral infections, cancer and genetic or metabolic diseases, comprises using gene chip and protein chip microarrays to identify specific DNA sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             sequence is a target region for an SDSO from an human cDNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 1; Length 19;
1.5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 19 BP; 5 A; 6 C; 7 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      / Match 4.2%; Score 19; DB Local Similarity 100.0%; Pred. No. 1.5 nes 19; Conservative 0; Mismatches
                                                                                                                                                                 Example 1; Page 19; 58pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        116 CAGCGGGGGGAAAAGCCTC 134
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 cagcegeedeaaaaeccrc 19
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ö Human; 88; SDSO; short double stranded oligonucleotide; cleavage site; viral infection; malignant tumour; genetic disease; metabolic disease; gene chip; protein chip; microarray; gene drug; Dermogene; Lungene; Hepatogene; Leukogene; Lymphogene; Prostogene; Breastogene; Human telomerase RNA gene, SDSO target region #6.

1 CCAGCCCCCGAACCCCGCC 19

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Braintumogene; Skin-whitogene; short interfering RNA; siRNA; cancer;
                              16-SEP-2002; 2002US-00016490.
                                     16-SEP-2002; 2002US-00016490.
                                                                          specific DNA sequences.
                                                         WPI; 2004-355427/33.
   RNA interference.
                 US2004072769-A1.
                                            (YINJ/) YIN J Q.
          Homo sapiens
                        15-APR-2004.
                                                   (in JQ)
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This invention relates to a novel method concerning prognostic markers associated with EGFR (epidermal growth factor receptor) positive cancer. Specifically, it refers to a gene expression profiling method that can provide a prediction as to whether a patient is likely to respond well to treatment with an EGFR inhibitor. The present invention describes the creatment with an EGFR inhibitor. The present invention describes the creatment with an EGFR inhibitor. The present invention describes the contractive analysis of the expression level of the RNA transcript of at CRP17, LAMC2 or their products thereof. It further provides a cDNA microarray containing named genes that represent prognostic transcripts which are useful for determining whether a patient diagnosed with an EGFR cappessing head or neck cancer or colon cancer exhibits elevated or decreased expression levels of these genes compared to normal. As such, these methods are also useful for prognosing or predicting the likelihood or cancer-attributable death or prognession, including recurrence and metastatic spread of a neoplastic disease, as well as drug resistance.

This oligonucleotide sequence is a PCR primer used to amplify a human PCR amplication of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Prognosing a patient with EGFR-expressing colon cancer comprises subjecting a sample comprising EGFR-expressing cancer cells to quantitative analysis of the expression level of the RNA transcript of at
                                                                                                                                                                                                                                              PCR primer to amplify a human cancer prognostic marker DNA SeqID 343.
                                                                                                                                                                                                                                                                                                     human; primer; PCR; prognostic marker; BGFR; epidermal growth factor receptor; cancer; gene expression profiling; microarray; head and neck cancer; colon cancer; metastatic spread;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Shak S, Baselga J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                claim 54; SEQ ID NO 343; 113pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          VALL-) VALL HEBRON UNIV HOSPITAL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15-NOV-2002; 2002US-0427090P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     14-NOV-2003; 2003WO-US036777.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      east one gene e.g., CD44v3.
                                                                ADP27906 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (GENO-) GENOMIC HEALTH INC
                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                      neoplastic disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Baker JB, Cronin MT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2004-420643/39.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO2004046386-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                        26-AUG-2004
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                                                                                                                             ADP27906;
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Matches
RESULT 2
ADP27906
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to screening, identifying or predicting, and assembling 19-25 at double-stranded oligonucleotides (termed short double stranded oligonucleotides, SDSO) as active pharmaceutical compositions cranded oligonucleotides, SDSO) as active pharmaceutical compositions for the treatment of viral infections, malignant tumours, and genetic and metabolic diseases, comprising screening and identifying a specific DNA sequence in an abnormal gene encoding a protein with gene chip and croarrays. The above method comprises screening the disease-causing genes, over-expressing in cells and/or tissues, with the composition of the part of the protein of plant protein chip microarrays. The above method comprises screening the disease-causing genes, over-expressing in cells and/or tissues, with the assistance of computer and specific DNA sequence within the assistance of computer and specific DNA composition of an RNA molectule and making sure that selected sequence is not localised within the stem-loop of target mRNA with any related software, long or 5-u(I)(TCCG-3' sepecial pattern complementary to at software. Also included are pharmaceutical compositions of general collections are pharmaceutical compositions of general collections. Braintungene and Skin-whitegene included are pharmaceutical compositions of general pattern for specific 19-25 in daRNA, 19-25 in sRNA-cDNA, 19-25 in daRNA and or more pharmaceutical carriers, one or more pecific call-targeting proteins and other active agents and additional capacing or not and a simplified method for pracial pattern that can be localised in any pecific and efficacious small double-stranded oligonucleotide sequence evaluating the specific and efficacious small double-stranded oligonucleotide sequence evaluating the specific and efficacious small addition, malignant tumours, general pattern that can be localided and sele
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               expression of genes in cells or tissues of humans or animals, such as viral infections, cancer, or genetic or metabolic diseases. The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Designing and selecting short double-stranded oligonucleotides for treating viral infections, cancer and genetic or metabolic diseases, comprises using gene chip and protein chip microarrays to identify
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   viral infections, cancer, or genetic or metabolic diseases.
sequence is a target region for an SDSO from an human cDNA.
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Pred. No. 1.5e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 1, Page 19; 58pp; English.
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Best Local Similarity 100.
Matches 19; Conservative
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                                                                        Gaps
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0
                                 . Match 4.2%; Score 19; DB 1; Length 19; Local Similarity 100.0%; Pred. No. 1.5e+02; es 19; Conservative 0; Mismatches 0; Indels
Sequence 19 BP; 7 A; 3 C; 8 G; 1 T; 0 U; 0 Other;
                                                                                                              371 AAGAGGAACGGAGCGAGTC 389
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Gaps

; 0

0; Indels

230 CCAGCCCCGAACCCCGCC 248

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'note = "2',4'-oxyethylene linkage in the sugar residues"
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/note= "2',4'-oxyethylene linkage in the sugar residues"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Stable 2',5'-oligoadenylic acid analogs containing natural and modified nucleic acid units as well as unusual phosphate groups with excellent activity particularly antitumor, applicable in cancer or antiviral
                                                                                                                               Cytostatic; virucide; 2'; 5'-oligoadenylic acid analog; antitumour;
                                                                                2',5'-oligoadenylic acid analog related oligonucleotide #4
                                                                                                                                                                                                                                                                                                                                                                                                              /mod_base= OTHER
/note= "phosphorothioate backbone"
15. 19
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/mod_base= OTHER
/note= "C-hydroxyethyl_phosphate"
                                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                                                                                                                                                                      mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19-NOV-2002; 2002JP-00334731.
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*tag= a
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                                                                                                                                                           antiviral; cancer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Morita K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2004-460494/43.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
nes 19; Conserv
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modified base
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                                   26-AUG-2004
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                                                                                                                                                                                                     Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This invention relates to a novel method concerning prognostic markers associated with EGFR (epidermal growth factor receptor) positive cancer. Specifically, it refers to a gene expression profiling method that can provide a prediction as to whether a patient is likely to respond well to treatment with an EGFR inhibitor. The present invention describes the quantitative analysis of the expression level of the RNA transcript of least one gene selected from the group of CD44v6, DBS, GROl, KRT17, LAMC2 or their products thereof. It further provides a cDNA microarray containing named genes that represent prognostic transcripts which are useful for determining whether a patient diagnosed with an EGFR expressing head or neck cancer or colon cancer exhibits elevated or decreased expression levels of these genes compared to normal. As such, these methods are also useful for prognessing or predicting the likelihood of cancer attributable death or progression, including recurrence and metastatic spread of a neoplastic disease, as well as drug resistance. This oligonucleotide sequence is a PCR primer used to amplify a human PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Prognosing a patient with EGFR-expressing colon cancer comprises subjecting a sample comprising EGFR-expressing cancer cells to quantitative analysis of the expression level of the RNA transcript of at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          amplicon DNA sequence used as a prognostic cancer marker, given in an exemplification of the invention.
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                                                                                                                                                                               PCR primer to amplify a human cancer prognostic marker DNA SeqID 344.
                                                                                                                                                                                                                       human; primer; PCR; prognostic marker; EGFR; expression profiling; epidermal growth factor receptor; cancer; gene expression profiling; microarray; head and neck cancer; colon cancer; metastatic spread;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
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Pred. No. 1.5e+02;
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100.0%; Pred. No. 1...
'... 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   54; SEQ ID NO 344; 113pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (GENO-) GENOMIC HEALTH INC.
(VALL-) VALL HEBRON UNIV HOSPITAL.
ADP27907/C

ADP27907 standard; DNA; 19 BP. XX
AC ADP27907;
XX
DE CAUG-2004 (first entry)
XX
DE FCR primer to amplify a human can
XX
W human; primer; PCR; prognostic m
W microarray; head and neck cancer
XX
M microarray; head and neck cancer
XX
M mocplastic disease; ss.
XX
Homo sapiens.
XX
Homo sapiens.
XX
HOMO-2003; 2002US-0427090P.
XX
M W2004046386-Al.
XX
H3-NOV-2002; 2002US-0427090P.
XX
M H3-NOV-2003; 2003WO-US036777.
XX
H3-NOV-2003; 2003WO-US036777.
XX
H3-NOV-2004; 2002US-0427090P.
XX
M H2: 2004-420643/39.
XX
GENO-) GENOMIC HEALTH INC.
XX
MPI; 2004-420643/39.
XX
GENO-) GENOMIC HEACH ECFR (epidermal)
XX
Claim S4; SEQ ID NO 344; 113pp; J
XX
Claim S4; SEQ ID NO 344; 113pp; J
XX
Claim S4; SEQ ID NO 344; 113pp; J
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Claim S4; SEQ ID NO 344; 113pp; J
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Claim S4; SEQ ID NO 344; 113pp; J
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Claim S4; SEQ ID NO 344; 113pp; J
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Claim S4; SEQ ID NO 344; 113pp; J
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Claim S4; SEQ ID NO 344; 113pp; J
XX
Claim S4; SEQ ID NO 344; 113
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                                   The present invention relates to novel 2',5'-oligoadenylic acid analogs and their pharmacologically- acceptable salts. The analogs are stable with superior antitumour and antiviral activity and so are useful in cancer or antiviral therapy e.g. as antisense drugs. The present sequence was used to illustrate the invention.
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                                                                                                                                                 Sequence 19 BP; 6 A; 5 C; 8 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                      Score 19; DB 1; Le
Pred. No. 1.5e+02;
                                                                                                                                                                                                                           0; Mismatches
Disclosure; Page 100; 220pp; Japanese.
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100.0%;
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                                                                                                                                                                                                                             Conservative
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ADP87879/C
ID ADP878
XX
AC ADP878
XX
DT 26-AUG
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ADP87877 standard; DNA; 19 BP.

RESULT 233
ADP87877/C
ID ADP878
XX
AC ADP878

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ADP87877

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Cytostatic; virucide; 2'; 5'-oligoadenylic acid analog; antitumour;
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 101; 220pp; Japanese.
                                                         ocation/Qualifiers
                                                                                                                                                                                                             /mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             76 GIGCTITIGCTCCCCGCGC 94
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Best Local Similarity 100. Matches 19; Conservative
                                                                                                                                                .19
                                                             antiviral; cancer; ss.
                                                                                                                                                                                                                                                                                                                          (SANY ) SANKYO CO LTD
                                                                                                                                                                                                                                                                                                                                             Koizumi M, Morita K;
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                                                           Key
modified_base
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                                                                                                                                                                                                                                                              03-JUN-2004
                                       Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                    therapy
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                                                                                                                                        /*tag= a
/mod_base= OTHER
/note= "2',4'-oxyethylene linkage in the sugar residues"
                                                                                                                                                                                 /*tag= c
/mod_base= OTHER
/note= "2',4'-oxyethylene linkage in the sugar residues"
                                                                                                                                                                                                                                                                                                                                                                                                        Stable 2',5'-oligoadenylic acid analogs containing natural and modified nucleic acid units as well as unusual phosphate groups with excellent activity particularly antitumor, applicable in cancer or antiviral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                             Cytostatic; virucide; 2'; 5'-oligoadenylic acid analog; antitumour;
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2',5'-oligoadenylic acid analog related oligonucleotide #7.
          2',5'-oligoadenylic acid analog related oligonucleotide #6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4.2%; Score 19; DB 1; Length 19;
100.0%; Pred. No. 1.5e+02;
ive 0; Mismatches 0; Indels
                                                                                                                      'note= "phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 19 BP; 6 A; 5 C; 8 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                            /note= "C-hydroxyethyl phosphate"
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 100-101; 220pp; Japanese.
                                                                            Location/Qualifiers
1. 19
/*tag= d
                                                                                                           base= OTHER
                                                                                                                                                                                                                                   base= OTHER
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Best Local Similarity luv...
Local 19; Conservative
                                         antiviral; cancer; ss
                                                                                                                                                                                                                                                                                                                                                (SANY ) SANKYO CO LTD
                                                                                                              /mod
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                                                                                 Key
modified_base
                                                                                                                                 modified base
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                                                             Synthetic.
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ID ADP8
XX ADP8
XX DT 26-A
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                                                                                                    *tag= a
/mod_base= OTHER
/note= "2',4'-oxyethylene linkage in the sugar residues"
                                                                                                                                                                                                                                                    /*tag= c
|mod_base= OTHER
|note= "2′,4′-oxyethylene linkage in the sugar residues"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Stable 2',5'-oligoadenylic acid analogs containing natural and modified nucleic acid units as well as unusual phosphate groups with excellent activity particularly antitumor, applicable in cancer or antiviral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4.2%; Score 19; DB 1; Length 19; 100.0%; Pred. No. 1.5e+02; tive 0; Mismatches 0; Indels
/mod_base= OTHER
/note= "phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note= "C-hydroxyethyl phosphate"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 19 BP; 6 A; 5 C; 8 G; 0 T; 0 U; 0 Other;
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misc_feature
                                     misc_feature
                                                                        WO9847911-A1
                                                                                                               13-APR-1998;
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03-FEB-1998;
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                                                                                                                                                                                                                                                                                                                         The present invention relates to novel 2',5'-oligoadenylic acid analogs and their pharmacologically- acceptable salts. The analogs are stable with supervior antitumour and antiviral activity and so are useful in cancer or antiviral therapy e.g. as antisense drugs. The present sequence was used to illustrate the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; telomerase; hTR; activator-antisense complex; malignant; enzyme; cleave; brain; tumour malignant glioma; breast tumour; renal cell cancer; melanoma; prostate cancer; leukemia; polycthemia vera; myeloma; sarcoma; Hodgkin; slymphoma; Maldenstrom s macroglobulinemia; heavy chain disease; carcinoma; chemotherapeutic; antisense; ss.
                                                               note- "2',4'-oxyethylene linkage in the sugar residues"
                                                                                                                                                                                                                                                        Stable 2',5'-oligoadenylic acid analogs containing natural and modified nucleic acid units as well as unusual phosphate groups with excellent activity particularly antitumor, applicable in cancer or antiviral
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Oligo contained activator-antisense complex spA12-anti-hTR.
                                                                                                                                                                                                                                                                                                                                                                                                    4.2%; Score 19; DB 1; Length 19; 100.0%; Pred. No. 1.5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                       0; Indels
                                                                                /ttag= b
/mod_base= OTHER
/note= "C-hydroxyethyl phosphate"
                                                                                                                                                                                                                                                                                                                                                                                  Sequence 19 BP; 6 A; 5 C; 8 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Prea. w..
                                                                                                                                                                                                                                                                                                        Disclosure; Page 100; 220pp; Japanese.
                           Location/Qualifiers
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                                               *tag= a
'mod base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                           76 GIGCITITGCICCCGCGC 94
                                                                                                                                                                                                                                                                                                                                                                                                                                                      GIGCTITIGCICCCCGCGC 1
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                                                                                                                                                            19-NOV-2003; 2003WO-JP014748
                                                                                                                                                                               19-NOV-2002; 2002JP-00334731
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1es 19; Conservative
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                                                                                                                                                                                                 (SANY ) SANKYO CO LTD
                                                                                                                                                                                                                    Koizumi M, Morita K;
                                                                                                                                                                                                                                        WPI; 2004-460494/43.
                                                                                                                       WO2004046161-A1
                                   modified base
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modified_base
                                                                         modified_base
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                                                                                                                                          03-JUN-2004.
         Synthetic.
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This represents an antisense oligonuclectide to the RNA component of human telomerase (hTR) comprised in the. The invention relates to an activator-antisense complex that comprises: (a) an antisense oligo, complementary to a 12-25 nuclectide portion of the RNA component of hTR, with a hydroxyl moiety at the first end; and (b) a linker attached to the linker attached to the cirst end, and (c) an activator of RNase L attached to the linker. The activator-antisense complex may be used for inhibiting the growth of a telomerase-expressing malignant cell or tumour. The complex is used to specifically cleave the ribonuclectide portion of a telomerase enzyme. The complex inhibits growth of telomerase expressing malignant cells from real norm, and prostate cancer. Many other malignancies and related disorders, may be treated including various acute and chronic leukemias, polycthemia vera, Hodgkin's and non-Hodgkin's lymphomas, multiple myelome, including numerous sarcomas and carcinomas. The complex is preferably administered in combination with a chemotherapeutic agent, complexin, actinomycin D, or necestinostetin. The present sequence is an example of a modified antisense oligo comprised in an activator-antisense
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New RNase L activator-telomerase antisense complex - useful to inhibit telomerase activity in telomerase-expressing malignancies.
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                                                                                                                                                                      /*tag= c
/note= "nucleotide in reverse orientation 3'-5'"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Torrence PF;
                                                                                                 /note= "3'-3' internucleotide linkage"
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"Sp5'A(2'p5'A)11-Bu2"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Li G,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example; Page 53; 81pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (CLEV-) CLEVELAND CLINIC FOUND. (USSH ) US NAT INST OF HEALTH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP
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98US-00018125.
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nes 19; Conservative
                               19. .20
/*tag= }
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                                                                                                                                                                   /*tag=
      'note=
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19 GIGCTTTIGCICCCGCGC 1

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melanoma; prostate cancer; leukemia; polycthemia vera; myeloma; sarcoma;
Hodgkin's lymphoma; Waldenstrom's macroglobulinemia; heavy chain disease;
carcinoma; chemotherapeutic; antisense; ss.
cleave; brain; tumour malignant glioma; breast tumour; renal cell cancer;
                                                                                                /note= "antisense oligo sequence claimed in claim 4"
                                                                                                                                                                      /*tag= d
/note= "nucleotide in reverse orientation 3'-5'"
                                                                                                                                                                                                                                                                                                                                                    telomerase activity in telomerase-expressing malignancies
                                                                                                                                           /*tag= c
note= "3'-3' internucleotide linkage"
                                                                                                                 '*tag= a
'note= "Sp5'A(2'p5'A)3-Bu2"
                                                                                                                                                                                                                                                                                                        Li G,
                                                                                                                                                                                                                                                                                                                                            New RNase L activator-telomerase antisense
                                                                                                                                                                                                                                                                                                        Silverman RH, Kondo S, Cowell JK,
                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                      Claim 4; Page 45; 81pp; English.
                                                                                                                                                                                                                                                                               CLEVELAND CLINIC FOUND
                                                                                                                                                                                                                                                            98US-00018125.
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                                                                                .19
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                                                                                                       modified_base
                                                     Homo sapiens
                                                                                                                                                            misc_feature
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                                                                             misc_feature
                                                                                                                                  misc_feature
                                                                                                                                                                                                                  29-OCT-1998
                                            Synthetic.
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this represents an antisense oligonucleotide to the RNA component of human telomerase (hTR) comprised in the. The invention relates to an activator-antisense complex that comprises: (a) an antisense oligo, activator-antisense complex that comprises: (a) an antisense oligo, with a hydroxyl moiety at the first end; and (b) a linker attached to the linker attached to the linker attached to the linker. The activator-antisense complex may be used for inhibiting the growth of a telomerase expressing malignant cell or tumour. The complex is used to specifically cleave the ribonucleotide portion of a telomerase enzyme. The complex inhibits growth of telomerase expressing malignant glioma, breast tumour, renal cell cancer. The complex inhibits growth of telomerase expressing malignant glioma, breast tumour, renal cell cancer. The complex inhibits drown to telomerase expressing malignant glioma, breast tumour, renal cell cancer. Collicating various acute and chronic leukemias, polycthemia vera, Hodgkin's and non-Hodgkin's lymphomas, multiple tumours, including numerous sarcomas and carcinomas. The complex is preferably administered in combination with a chemotherapeutic agent, bleomychin, actinomycin D, or neocarzinostatin. The present sequence is an example of a modified antisense oligo comprised in an activator-antisense
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Claim 11; Page 13; 52pp; English.
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100.0%; Pred. No. 1.6e+02;
tive 0; Mismatches 0; Indels
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The present sequence is that of a short interfering RNA, denoted hTR#2 siRNA, a 19 bp sequence centred in the 26 bp loop in the longest single-stranded region of human telemerase RNA. The siRNA comprises sense and antisense nucleic acids that are complementary to each other except for 2 thymidine deoxynucleotides at both 3' overhangs. A nucleic acid comprising the double-stranded region of this siRNA is specifically claimed. The invention relates to the discovery that double-stranded interfering RNAs which target telomerase RNA, or mRNA encoding telomerase interfering RNAs which target telomerase RNA, or mRNA encoding telomerase activity. Inhibition of telomerase in cancer cells leads to telomerase activity end-to-end chromosomal fusion, and apoptosis. Interference of shortening, end-to-end chromosomal fusion, and apoptosis. Interference of contraception or sterilisation, for immunosuppression, for treatment of contraception or sterilisation, for immunosuppression, for treatment of yeast, parasite and fungal infections, and in antiinflammatory therapies. As telomerase is active in a limited number of cell types, e.g. tumour cells, germaine cells, certain stem cells of the haematopoietic system, T and B cells, sun-damaged skih, and proliferative cervix, most normal cells are not affected by telomerase RNA interference therapy
                                                                                                                                                                                                                                                                                                                                                                                                                               /*tag= a //tabel= Sticky end //tabel= Sticky end //tabel= "the 3' end of the complementary strand overhangs the s' end of this strand by the sequence 5'-IT-3'"
                                                                                                                                                                                                                                                                                                                                                                          "double-stranded region, specifically referred to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel nucleic acid encoding or comprising interfering RNAs which target telomerase RNA, useful for inhibiting telomerase activity for treating cancer, infertility and disorders of the immune system.
                                                                                                                                                                                          RNA interference; short interfering RNA; siRNA; telomerase; cancer; tumour; cytostatic; contraceptive; immunosuppressive; antiinfertility; fungicide; antiparasitic; antiinflammatory; human; gene therapy; ds.
                                                                                                                                                     Short interfering RNA hTR#2 siRNA, targets telomerase RNA.
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/label= Sticky_end
                                      BP.
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20-FEB-2002; 2002US-0359196P.
22-MAY-2002; 2002US-0383195P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16-OCT-2002; 2002WO-US033146.
                                                                                                                                                                                                                                                                                                                                                                                            in Claim 11"
                                    ACC57540 standard; DNA; 21
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*tag= b
                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                              note=
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                                                                                                                                                                                                                                                                                                                  Key
misc_feature
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                                                                           ACC57540;
RESULT 239
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complex - useful to inhibit

Torrence PF;

76 GIGCITITGCICCCGCGC 94

Local Similarity 100. Les 19; Conservative

Matches

Query Match

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WO2003070742-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   modified_base
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29-AUG-2002;
05-SEP-2002;
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06-JUN-2002;
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                                                                                                                                Query Match
                                                                                                                                                                                                                        RESULT 241
ADF93815/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present sequence is that of a short interfering RNA, denoted hTR#1 siRNA, which targets the template region (see ACG57538) of human telomerase RNA. The siRNA comprises sense and antisense nucleic acids that are complementary to each other except for 2 thymidine deoxynucleotides at both 3' overhangs. A nucleic acid comprising the duble-stranded region of this siRNA is specifically claimed. The invention relates to the discovery that double-stranded interfering RNAs which target telomerase RNA, or mRNA encoding telomerase reverse
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/label= Sticky_end
/note= "the 3' end of the complementary strand overhangs
the 5' end of this strand by the sequence 5'-TT-3'"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel nucleic acid encoding or comprising interfering RNAs which target telomerase RNA, useful for inhibiting telomerase activity for treating cancer, infertility and disorders of the immune system.
                                                                                                                                                                                                                  RNA interference; short interfering RNA; siRNA; telomerase; cancer; tumour; cytostatic; contraceptive; immunosuppressive; antiinfertility; fungicide; antiparasitic; antiinflammatory; human; gene therapy; DNA-RNA hybrid; ds.
                                                                                                                                                                                                                                                                                                                         /note= "double-stranded region, specifically referred
in Claim 10"
                                                 Gaps
                                                 ö
                                                                                                                                                                                                    Short interfering RNA hTR#1 siRNA, targets telomerase RNA
                            Score 19; DB 1; Length 21; Pred. No. 1.7e+02;
                                                0; Indels
          Sequence 21 BP; 2 A; 8 C; 6 G; 5 T; 0 U; 0 Other;
                                  100.0%; Prea. ....
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/label= Sticky_end
                                                                   270 GGCTTCTCCGGAGGCACCC 288
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                                                                                       1 GGCTTCTCCGGAGGCACCC 19
                                                                                                                                        ACC57539 standard; DNA; 21 BP
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2002US-0383195P.
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                             4.28;
                                                                                                                                                                                (first entry)
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                                      Best Local Similarity 100.
Matches 19; Conservative
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                             Query Match
                                                                                                                     RESULT 240
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transcriptase (TERT), are capable of inhibiting telomerase activity. Inhibition of telomerase in cancer cells leads to telomera shortening, end-to-end chromosomal fusion, and apoptosis. Interference of telomerase activity can also be used for treatment of infertility, for contraception or sterilastion, for immunosuppression, for treatment of of yeast, parasite and fungal infections, and in antiinflammatory therapies. As telomerase is active in a limited number of cell types, e.g. tumour cells, germline cells, certain stem cells of the hematopojoitic system, T and B cells, sun-damaged skin, and proliferative cervix, most normal cells are not affected by telomerase RNA interference therapy
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                                                                                                                                                                                                                                                                                                                                                                                     Sequence 21 BP; 5 A; 5 C; 3 G; 2 T; 6 U; 0 Other;
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/mod base= OTHER
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2002US-0363124P.
2002US-0386782P.
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2002US-0408378P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity 68.4
nes 13; Conservative
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and matisense regions, or alternatively are assembled from a sense oligonuclectide and an interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short interfering RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonuclectides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA, conjugates candor complexes of siNA, and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and cranded the confictions of the telomerase genes in cells, tissue companisms (e.g., by ex vivo gene therapy), or in grafts and confictions of the treatment of a variety of conditions. They may be conditions or organisms (e.g., by expense).
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                                                                                                                                                                                                                                                                                                                                                                                                                                       used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siMs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nuclectide polymorphisms). The present sequence represents a chemically modified siRNA targeted to the human TERC mRNA
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telomerase gene useful e.g. for treatment and diagnosis of cancer.
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4.2%; Score 19; DD.
Best Local Similarity 100.0%; Pred. No. 1.7.
Best Local Similarity 0, Mismatches
                                              Example 3; SEQ ID NO 542; 145pp; English.
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense cregions, or alternatively are assembled from a sense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double stranded RNA, micro-RNA (miRNA) and short hairpin RNA (shRNA). The siNAs can be unmodified or chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised.

CC can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised.

CC relates to kits for the in vitro or in vitro of in vitro of in the cxpress sinA. The sinAs are cand/or complexes of sinA; and vectors that express sinA. The sinAs are explaints or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be conditioned in transplant rejection, or autoimmune or age-related diseases, sin multiple sclerosis, lupus erythematosus, AlDS, macular degeneration, and rheumatoid arthritis. The sinAs are also useful for drug
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        genetic engineering, pharmacogenomics, studying gene function, and gene napping (e.g., of single nucleotide polymorphisms). The present sequence represents a chemically modified siRNA targeted to the human TERC mRNA
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100.0%; Pred. No. 1.7e+02;
ive 0; Mismatches 0; Indels
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15-JAN-2003; 2003US-0440129P.
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"...hes 19; Conservative
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and mantisense regions, or alternatively are assembled from a sense oligonuclectide and an interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short interfering RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonuclectides, and can be chemically synthesised, can contain deoxyribonuclectides, and can be chemically synthesised, capressed from a vector or enzymatically synthesised. The invention also relates to Kits for the in vitro or in vivo delivery of siNA, conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are suplants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be creating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, shill ulcers and rheumacoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence
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/note= "Ribothymidines. Also, the internucleotide linkage
is phosphorothioate"
Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiartits; antiartits; antiartits; antiartits; antiartits; antiartits; antiartits; antiartits; santiartits; santiartitaring nucleic acid; sinA; short interfering NNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERT;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           29-AUG-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20-FEB-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17-JUL-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          28-AUG-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic
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                                                                                                                                                                                                                                                                                                                                                      Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiartitic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERT; DNA-RNA hybrid; se.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense
represents a chemically modified siRNA targeted to the human TERC mRNA
                                                                                                     Gарв
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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                                                                        4.2%; Score 19; DB 1; Length 21;
                                                                                                  Indele
                                                                                                                                                                                                                                                                                                                             Human TERC chemically modified siRNA, SEQ ID 539.
                                           Seguence 21 BP; 7 A; 2 C; 6 G; 2 T; 4 U; 0 Other;
                                                                                       1.7e+02;
                                                                                                     0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 3; SEQ ID NO 539; 145pp; English.
                                                                                         Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note= "Ribothymidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                   146 TTCCACCGTTCATTCTAGA 164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                base= OTHER
                                                                                                                                                    TTCCACCGTTCATTCTAGA 1
                                                                                                                                                                                                                                       ADF93812 standard; RNA; 21 BP
                                                                                      100.08;
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2002US-0406784P.
2002US-0408378P.
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2002US-0386782P
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                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                     19; Conservative
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                                                                                      Best Local Similarity
Matches 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO2003070742-A1
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11-MAR-2002;
06-JUN-2002;
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05-SEP-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28-AUG-2003
                  transcript.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                             19
                                                                                                                                                                                                                                                                   ADF93812;
                                                                          Query Match
                                                                                                                                                                                                           244
                                                                                                                                                                                                          RESULT
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regions, or alternatively are assembled from a sense oligonucleotide and antisense oligonucleotide. Specifically, the sinAs include short interfering RNA (sinKA), double-stranded RNA, micro-NRA (miRNA) and short hairpin RNA (shRNA). The sinAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of sinA; conjugates and/or complexes of sinA; and vectors that express sinA. The sinAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants of organisms (e.g., by ex vivo gene therapy), or in grafts and transplants of organisms (e.g., by ex vivo gene therapy), or in grafts and transplant rejection, or autoimmune or age-related diseases, protozoal), transplant rejection, or autoimmune or age-related diseases, c.g. multiple sclerosis, lupus erythematosus, AlDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence mapping (e.g., of single nucleotide polymorphisms). The present sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                4.2%; Score 19; DB 1; Length 21; 73.7%; Pred. No. 1.7e+02; ive 5; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                Sequence 21 BP; 4 A; 7 C; 3 G; 2 T; 5 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human TERC chemically modified siRNA, SEQ ID 552.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           148 CCACCGTTCATTCTAGAGC 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ВР
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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ADF93825/c
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New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.

05-SEP-2002; 2002US-0408378P. 09-SEP-2002; 2002US-0409293P. 15-JAN-2003; 2003US-0440129P.

(RIBO-) RIBOZYME PHARM INC

Beigelman L;

Mcswiggen J,

WPI; 2003-689777/65.

2002US-0396782P. 2002US-0396600P. 2002US-0406784P. 2002US-0363124P

06-JUN-2002; 17-JUL-2002; 29-AUG-2002;

11-MAR-2002;

Example 3; SEQ ID NO 552; 145pp; English.

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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense or regions, or alternatively are assembled from a sense onigonucleotide and antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short compressed from a vector or enzymatically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, can dor complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue capants for the treatment of a variety of conditions. They may be transplants for the treatment of a variety of conditions. They may be conditions for treating cancer, respenses infectious diseases (specifically proteozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, ALDS, macular degeneration, stin ulcers and rheumatoid arthritis. The siNAs are also useful for drug cortening, diagnosis, lupus erythematosus, ALDS, macular degeneration, canetic engineering, pharmacogenomics, studying gene function, and gene corporation. The present sequence contained a chemically modified siRNA targeted to the human TERC mRNA
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100.0%; Pred. No. 1.7e+02;
ive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 21 BP; 5 A; 7 C; 3 G; 2 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      300 GAAGAGTTGGGCTCTGTCA 318
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19 GAAGAGTTGGGCTCTGTCA 1
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Best Local Similarity 100.
Matches 19; Conservative
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/note= "Ribothymidines. Also, the internucleotide linkage
is phosphorothioate"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; mixNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT;
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*tag= b
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modified_base

Homo sapiens.

Synthetic

modified_base

28-AUG-2003.

0;

Gaps

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ADF93832/
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     mapping (e.g., of single nucleotide polymorphisms). The present sequence represents a chemically modified siRNA targeted to the human TERC mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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100.0%; Pred. No. 1.7e+02;
ive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                 /note= "Ribothymidines"
                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                    *tag= a
mod_base= OTHER
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2002US-0406784P.
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2002US-0386782P
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DNA-RNA hybrid; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                               WO2003070742-A1.
                                                                                                                                                                                                                                   modified base
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29-AUG-2002;
                                                                                                                          Homo sapiens
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                                                                                 Synthetic
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antiennse regions, or alternatively are assembled from a sense oligonucleotide and an antistense oligonucleotide. Specifically, the siNAs include short interfering RNA (siNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siNA), double-stranded RNA, micro-RNA (miRNA) and short can contain deoxyribonucleotides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised, me invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                /*tag= a
/mod_base= OTHER
/note= "Ribothymidines. Also, the internucleotide linkage
is phosphorothloate"
                                                                                                                            Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERT;
                                                                                                                                                                                                                                                                                                                                                                                                                 /note= "Pyrimidine bases are 2'-deoxy-2'-fluoro and purines are Deoxy bases"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
                                                                                             Human TERC chemically modified siRNA, SEQ ID 559.
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                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                  /mod_base= OTHER
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2002US-0408378P.
2002US-0409293P.
2003US-0440129P.
ADF93832 standard; RNA; 21 BP
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2002US-0386782P.
2002US-0396600P.
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                                                              (first entry)
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/*tag=
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modified_base
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                                                                26-FEB-2004
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05-SEP-2002;
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                                                                                                                                                                                                                                                                                                   Synthetic
                                ADF93832;
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Gaps

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300 GAAGAGTIGGGCTCTGTCA 318

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Conservative

Local Similarity les 19; Conserv

Matches

19 GAAGAGTTGGGCTCTGTCA 1

US09436060A-16.rng.sl

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used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenceis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple solarons in Jupus erythematocusus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siMAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single mucleotide polymorphisms). The present sequence represents a chemically modified siRNA targeted to the human TERC mRNA
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                                                                                                                                                                                                                                                  Length 21;
                                                                                                                                                                                                                                                                                   0; Indels
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100.0%; Pred. No. 1.7e+02;
iive 0; Mismatches 0;
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/mod_base= OTHER
/note= "Ribothymidines"
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                                                                                                                                                                                                                                                                                                                     CCACCGTTCATTCTAGAGC 166
                                                                                                                                                                                                                                                                                                                                        CCACCGTTCATTCTAGAGC 1
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11-MAX-2002; 2002US-0363124P.
06-UJN-2002; 2002US-0386782P.
17-UJL-2002; 2002US-0406784P.
29-AUG-2002; 2002US-0406784P.
09-SEP-2002; 2002US-0406778P.
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                                                                                                                                                                                                                                                                    Local Similarity 100.
nes 19; Conservative
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modified_base
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Homo sapiens
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                                                                                                                                                                                                                                                                                                                     148
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                                                                                                                                                                                                                                                                                      Matches
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WPI; 2003-689777/65.

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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and an attisense oligonucleotide. Specifically, the siNAs include short hairpin RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, capted to kits for the in vitro or in vivo delivery of siNA, conjugates and/or complexes of siNA, and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue to explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug
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                 New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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4.2%; Score 19; DB 1; Length 21;
Best Local Similarity 63.2%; Pred. No. 1.7e+02;
Matches 12; Conservative 7; Mismatches 0; Indels
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                                                                                         Example 3; SEQ ID NO 538; 145pp; English.
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/mod_base= OTHER
/note= "Ribothymidines"
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Homo sapiens
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06-JUN-2002;
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29-AUG-2002;
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                                                                                                Synthetic
                                                                                                                                           Κeγ
 The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and antisense constant sense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised of can contain deoxyribonucleotides, and can be chemically synthesised. CC expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express sink. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically proteozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AlDS, macular degeneration, skin ulcers and rheumatoid arthritis. The sinAs are also useful for drug screening, disgenses the canger deentification, and gene containing, disgenses the plymorphisms). The present sequence containing its end of a single mucleocide polymorphisms). The present sequence contains.
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                                                                                                                                                                                                                                                                   New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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Pred. No. 1.7e+02;
5; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 21 BP; 4 A; 3 C; 7 G; 2 T; 5 U; 0 Other;
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17-JUL-2002; 2002US-039660P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0409378P.
09-SEP-2002; 2002US-0409293P.
                                                                     2002US-0363124P
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Best Local Similarity 73.7%;
Matches 14; Conservative
                          11-FEB-2003; 2003WO-US004088
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADF93833 standard; RNA; 21
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28-AUG-2003
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interference. The siNAs may or may not comprise sense and antibened interference. The siNAs may or may not comprise sense and antibened interference. The siNAs may or may not comprise ribonucleotides and may be devolved or single stranded. They further comprise sense and antibened regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (shRNA). The siNAs can be unmodified or chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA, conjugates and/or complexes of siNA, and vectors that express sink. The siNAs are used to modulate expression of the telomerase genes in cells, tissue used to modulate expression of the telomerase genes in cells, tissue used for treating cancer, restenosis, infectious diseases (specifically proteins or protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AlbS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, disgnossis, therappentic tazget identification and validation, and genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence constrains a chemically modified siRNA targeted to the human TERC mRNA
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/mod_base= OTHER
/note= "Ribothymidines. Also, the internucleotide linkage
is phosphorothioate"
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                                                   miRNA;
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20. .21
RNA interference; short interfering nucleic acid; siNA; miRNA short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; DNA-RNA hybrid; se.
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2002US-0386782P.
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2002US-0406784P.
2002US-0408378P.
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11-MAR-2002;
06-JUN-2002;
17-JUL-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 252
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADF93823/
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cytostatic, vasotropic, protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleacid; siNA; short interfering nucleacid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; DNA-RNA hybrid; ss.
                                                                                                                        Gaps
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                                                                                                                           ;
0
                                                            Length 21;
                                                                                                                        0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human TERC chemically modified siRNA, SEQ ID 543.
   BP; 5 A; 7 C; 3 G; 2 T; 4 U; 0 Other;
                                                         4.2%; Score 19; DB 1; Le
100.0%; Pred. No. 1.7e+02;
:ive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                        300 GAAGAGTTGGGCTCTGTCA 318
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   base= OTHER
                                                                                                                                                                                                                       GAAGAGTTGGGCTCTGTCA 1
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2002US-0363124P.
2002US-0386782P.
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29-MUG-2002; 2002US-0406784P-
05-SEP-2002; 2002US-0408378P-
09-SEP-2002; 2002US-0409293P-
15-JAN-2003; 2003US-0440129P-
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                                                                                                                                                                                                                                                                                                                                                                                                      ADF93816 standard; RNA; 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                            Local Similarity 100.
nes 19; Conservative
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20-FEB-2002;
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         Sequence 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADF93816;
                                                                                                                                                                                                                                                    19
                                                                  Query Match
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                                                                                                                                  Matches
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hairpin RNA (shRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits a vector or in vivo delivery of siNA; conjugates and/or complexes of siNA, and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antinheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; expension modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents a chemically modified siRNA targeted to the human TERC mRNA
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/note= "Pyrimidine bases are 2'-deoxy-2'-fluoro"
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100.0%; Pred. No. 1.7e+02;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human TERC chemically modified siRNA, SEQ ID 550.
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/mod base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                148 CCACCGTTCATTCTAGAGC 166
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2002US-0363124P.
2002US-0386782P.
2002US-0396600P.
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Matches 19; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO2003070742-A1
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modified_base
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                                                                                                                                                                                                                            The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonuclectide and an interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs and can be chemically synthesised, can contain deoxyribonuclectides, and can be chemically synthesised, can contain deoxyribonuclectides, and can be chemically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA, conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue caplants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants of organisms (e.g., by ex vivo gene therapy), or in grafts and transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus exthematosus, AlbS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and gene empired engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single mulcleotide polymorphisms). The present sequence mapping (e.g., of single mulcleotide polymorphisms). The present sequence correction and acceptance of the constant and targeted to the human TERC mRNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            double-stranded short interfering nucleic acid; siNA; antiparkinsonian; antiarteriosclerotic; neuroprotective; noctropic; antiparkinsonian; anticonvulsant; pulmonary disease; restenosis; atherosclerosis; alzheimer's; Parkinson's; epilepsy; dementia; huntington's; amyotrophic lateral sclerosis; gene therapy; ss; DNA-RNA hybrid; hTR.
                                                                                                                                                         New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4.2%; Score 19; DB 1; Length 21;
100.0%; Pred. No. 1.7e+02;
live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 21 BP; 7 A; 2 C; 6 G; 2 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sina DNA-RNA hybrid - SEQ ID 608.
                                                                                                                                                                                                       Example 3; SEQ ID NO 550; 145pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           146 TTCCACCGTTCATTCTAGA 164
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29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-040929P.
15-JAN-2003; 2003US-0440129P.
                                                                      (RIBO-) RIBOZYME PHARM INC.
                                                                                                   Beigelman L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
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les 19; Conservative
                                                                                                                              WPI; 2003-689777/65.
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                                                                                                   Mcswiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   hTR-targeted
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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The invention relates to a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene comprising one or more chemical modifications and each strand of the double-stranded siNA comprises about 21 nucleotides. The siNA of the invention demonstrates antiarteriosclerotic, neuroprotective, continuous antiparkinsonian and anticonvulsant activities and may be useful for down-regulating the expression of an endogenous mammalian target gene and therefore in the treatment of any disease or condition that responds to modulation of gene expression or activity in a cell, tissue or organism. The disease or condition may include pulmonary diseases such as restenosis, atherosclerosis, Alzheimer's disease, Parkinson's disease or may condition the sine and therefore in the constitution may include pulmonary diseases such as restenosis, atherosclerosis, Alzheimer's disease or mayotrophic lateral sclerosis. Furthermore, the sinA may be utilised for gene therapy applications. The current sequence is that of the sinA DNA-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New double-stranded short interfering nucleic acid molecule, useful for down-regulating the expression of an endogenous mammalian target gene or for treating diseases that respond to modulation of gene expression or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 19; DB 1; Length 21;
Pred. No. 1.7e+02;
0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                   Pavco P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 21 BP; 7 A; 2 C; 6 G; 2 T; 4 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                   Chowrira B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Thompson J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TICCACCGIICATICIAGA 164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TTCCACCGTTCATTCTAGA 1
                                                                                                                                                                                                                                                                                                                                                                      (SIRN-) SIRNA THERAPEUTICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity 100.0%; es 19; Conservation
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2002US-0386782P.
                                                                                                                                                                                                            2002US-0406784P.
                                                                                                                                                                                                                                                                                                         15-JAN-2003; 2003US-0440129P.
                                                         20-FEB-2003; 2003WO-US005028
                                                                                                                                                                                                                                                                            2002US-0409293P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                   Beigelman
                                                                                                                                                                                                                                                                                                                                                                                                      Mcswiggen J, Belyerm.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-731676/69.
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                                                                                                                                                                                                            29-AUG-2002;
                                                                                                                                                                                                                                                05-SEP-2002;
                                                                                                                                                                                                                                                                            09-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unidentified
                                                                                                                                                                                   36-JUN-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             26-FEB-2004
12-SEP-2003
                                                                                                                                                   11-MAR-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADG30043;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 254
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WO2003074654-A2
                      12-SEP-2003
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                                                                                                                                                                                                                                                                                                                                 (sinA) molecule that down-regulates expression of an endogenous mammalian target gene comprising one or more chemical modifications and each strand of the double-stranded sinA comprises about 21 nucleotides. The sinA of the invention demonstrates anitarteriosclerotic, neuroprotective, noctropic, antiparkinsonian and anticonvulsant activities and may be neefful for down-regulating the expression of an endogenous mammalian target gene and therefore in the treatment of any disease or condition that responds to modulation of gene expression or activity in a cell, rissue or organism. The disease or condition may include pulmonary diseases such as restenosis, atherosclerosis, Alzheimer's disease. Parkinson's disease, epilepsy, dementia, huntington's disease or agantism atherosclerosis, huntington's disease or amontorophic lateral sclerosis. Purthermore, the sinA may be utilised for gene therapy applications. The current sequence is that of the sinA DNA-RNA hybrid of the invention.
                                                                                                                                                                                                                                                                                                                         The invention relates to a double-stranded short interfering nucleic acid
                                                                                                                                                                                                                                      New double-stranded short interfering nucleic acid molecule, useful for down-regulating the expression of an endogenous mammalian target gene or for treating diseases that respond to modulation of gene expression or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       double-stranded short interfering nucleic acid; siNA; antiarteriosclerotic; neuroprotective; noctropic; antiparkinsonian; anticonvulsant; pulmonary disease; restenosis; atherosclerosis; alzheimer's; Parkinson's; epilepsy; dementia; huntington's; amyotrophic lateral sclerosis; gene therapy; ss; DNA-RNA hybrid; hTR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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0
                                                                                                                                                                               Fosnaugh K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 19; DB 1; Length 21;
Pred. No. 1.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Indels
                                                                                                                                                                               Pavco P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 21 BP; 5 A; 3 C; 7 G; 2 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 hTR-targeted sinA DNA-RNA hybrid - SEQ ID 610.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%; Prec. ...
                                                                                                                                                                                                                                                                                                   Example 24; SEQ ID NO 609; 593pp; English.
                                                                                                                                                                               Chowrira B,
                                                                                                                                                                          Beigelman L, Chowr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         148 CCACCGTTCATTCTAGAGC 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CCACCGTTCATTCTAGAGC 1
                                                                                                                                                       (SIRN-) SIRNA THERAPEUTICS INC
                                                                                2002US-0386782P.
2002US-0406784P.
2002US-0408378P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADG30044 standard; RNA; 21 BP
                                                         2002US-0358580P.
2002US-0363124P.
                                                                                                                    2002US-0409293P.
                                                                                                                                  15-JAN-2003; 2003US-0440129P.
                                   20-FEB-2003; 2003WO-US005028
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19; Conservative
                                                                                                                                                                  Mcswiggen J, Belywan N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
                                                                                                                                                                                                                 WPI; 2003-731676/69.
                                                                                           29-AUG-2002;
05-SEP-2002;
09-SEP-2002;
                                                                                 06-JUN-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          26-FEB-2004
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                                                                      11-MAR-2002;
            12-SEP-2003.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADG30044;
                                                                                                                                                                                                                                                                             activity.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 255
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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The invention relates to a double-stranded short interfering nucleic acid (siN) molecule that down-regulates expression of an endogenous mammalian target gene comprising one or more chemical modifications and each strand of the double-stranded siNA comprises about 21 nucleotides. The siNA of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New double-stranded short interfering nucleic acid molecule, useful for down-regulating the expression of an endogenous mammalian target gene or for treating diseases that respond to modulation of gene expression or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RNA interference; short interfering RNA; siRNA; telomerase; cancer; tumour; cytostatic; contraceptive; immunosuppressive; antiinfertility; fungicide; antiparasitic; antiinflammatory; human; gene therapy; DNA-RNA hybrid; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                    Fosnaugh K;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 19; DB 1; Length 21; Pred. No. 1.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                    Chowrira B, Pavco P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 21 BP; 5 A; 7 C; 3 G; 2 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; Pred. .v.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 24; SEQ ID NO 610; 593pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                          Beigelman L, Chowri
sman N, Thompson J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   300 GAAGAGTTGGGCTCTGTCA 318
                                                                                           11-MAR-2002; 2002US-0363124P.
06-JUN-2002; 2002US-0386782P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
09-SEP-2003; 2002US-040923P.
15-JAN-2003; 2003US-0440129P.
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                                                                                                                                                                                                                                                                                                                                  (SIRN-) SIRNA THERAPEUTICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP.
20-FEB-2003; 2003WO-US005028
                                                               2002US-0358580P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RNA hybrid of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               11-AUG-2003 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 100.
Matches 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                    Usman N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-731676/69.
                                                                                                                                                                                                                                                                                                                                                                                                    Mcswiggen J,
                                                               20-FEB-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                    Jamison S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                                                                                                                           Novel double-stranded short interfering RNA having sense and antisense unductaic acids which are complementary to each other and to target nucleic acid e.g., telomerase RNA or mRNA encoding telomerase reverse transcriptase.
           /*tag= b
/note= "double-stranded region, specifically referred to
                                                           /*tag= a
/label= Sticky_end
/note= "the 3' end of the complementary strand overhangs
the 5' end of this strand by the sequence 5'-TT-3'"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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100.0%; Pred. No. 1.7e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%; Preα. ....
rive 0; Mismatches
                                                                                                                        /*tag= c
/label= Sticky end
                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 11; Page 28; 37pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                270 GGCTTCTCCGGAGGCACCC 288
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ACC58027 standard; DNA; 21 BP
                                                                                                                                                                                                                                  22-OCT-2001; 2001US-0345326P.
20-FEB-2002; 2002US-0359196P.
22-MAY-2002; 2002US-0383195P.
                                                                                                                                                                                                          16-OCT-2002; 2002WO-US033065
                                  in Claim 11"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
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                                                                                                                                                                                                                                                                                   (UYRP ) UNIV ROCHESTER
                                                                                                                                                                                                                                                                                                                                     WPI; 2003-403336/38
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Matches 19; Conserv
                                                                                                                                                           WO2003035667-A2
                                                                                                                                                                                                                                22-OCT-2001;
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                                                                                                        misc_feature
 misc_feature
                                                 misc_feature
                                                                                                                                                                                   01-MAY-2003
                                                                                                                                                                                                                                                                                                           Rowley PT;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present sequence is that of a short interfering RNA, denoted hTR#1 siRNA, which targets the template region (see ACC57526) of human telenomerase RNA. The siRNA comprises sense and antisense nucleic acids that are complementary to each other except for 2 thymidine decoxput lectides at both 3 overhangs. A nucleic acid comprising the double-stranded region of this siRNA is specifically claimed. The invention relates to the discovery that double-stranded interfering RNAs which target telomerase RNA, or mRNA encoding telomerase reverse transcriptase (TERT), are capable of inhibiting telomerase activity. Inhibition of telomerase in cancer cells leads to telomerase activity end-to-end chromosomal fusion, and apoptosis. Interference of telomerase activity can also be used for treatment of infertility, for contraception or sterilisation, for imminosuppression, for treatment of season.
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                                                                                                                                                                                                                                                                                                                                                                                        /*tag= b
/note= "double-stranded region, specifically referred to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        overhangs
                                                                   RNA interference; short interfering RNA; siRNA; telomerase; cancer; tumour; cytostatic; contraceptive; immunosuppressive; antiinfertility; fungicide; antiparasitic; antiinflammatory; human; gene therapy; DNA-RNA hybrid; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /*tag= a
/label= Sticky_end
/note= "the 3' end of the complementary strand ove
the 5' end of this strand by the sequence 5'-TT-3'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
Short interfering RNA hTR#1 siRNA, targets telomerase RNA.
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                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
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/label= Sticky_end
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         42 TIGICIAACCCIAACTGAG 60
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20-FEB-2002; 2002US-0359196P.
22-MAY-2002; 2002US-0383195P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               in Claim 10"
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Best Local Similarity 68.41
Matches 13, Conservative
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*tag=
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Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA, expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to short interfering nucleic acids (siNA) which
                                                                                                                                                                                                                                                                                                                                                        /note= "Pyrimidine bases are 2'-deoxy-2'-fluoro and purines are Deoxy bases"
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/mod_base= OTHER
/note= "Inverted deoxy abasic nucleotide"
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/mod_base= OTHER
/note= "Inverted deoxy abasic nucleotide"
21. .22
                                                                                                      Human TERC chemically modified siRNA, SEQ ID 556.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              note= "Ribothymidines"
                                                                                                                                                                                                                                                                                                         Location/Qualifiers
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/mod_base= OTHER
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11-MAR-2002, 2002US-0363144P.
06-JUN-2002, 2002US-03660B2.
17-JUL-2002, 2002US-036600P.
29-AUG-2002, 2002US-0406784P.
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                          ADF93829 standard; RNA; 23
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                                                                             (first entry)
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                                                                                                                                                                                                                                        DNA-RNA hybrid; ss.
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                                                                                                                                                                                                                                                                                Homo sapiens
                                                                              26-FEB-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        09-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      28-AUG-2003
                                                                                                                                                                                                                                                                  Synthetic
                                                    ADF93829;
RESULT 25
ADF93829
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cc downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise sense and antisense consideratively are assembled from a sense and antisense regions, or alternatively are assembled from a sense only and antisense cregions, or alternatively are assembled from a sense only and antisense creating RNA (sinRNA), double-stranded RNA, micro-RNA (miKNA) and short hairpin RNA (shRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also crelates to kits for the in vitro or in vivo delivery of siNA, conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treatging cancer, restenosis, infectious diseases (specifically conditions) will unders and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, standard sequence mapping (e.g., of single mucleotide polymorphisms). The present sequence represents a chemically modified siRNA targeted to the human TERC mRNA
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/note= "Pyrimidine bases are 2'-deoxy-2'-fluoro and
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/note= "Inverted deoxy abasic nucleotide"
/note= "Inverted deoxy abasic nucleotide"
/*tag= c
/*mod_base= OTHER
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/note= "Ribothymidines"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP.
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Best Local Similarity 73.7%;
Matches 14; Conservative
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modified base

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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (shRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue captains (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically proteozal), transplant rejection, or autoimmune or age-related diseases, proteozal), transplant rejection, or autoimmune or age-related diseases, proteozoal), transplant rejection, or autoimmune or age-related diseases, computing all and relation and validation, cancer, near and cancer or stransplants for the sinas are also useful for drug cancer, cancerning, diagnosis, therapeleoget identification and validation, cancer or cancerning, diagnosis, therapeleoget identification and validation, cancer or cancerning, diagnosis, therapeleoget identification and validation, cancer or cancerning, diagnosis, therapeleoget identification and validation, and general or cancer or cancerning, diagnosis, therapeleoget identification and validation, and general cancer or cancernance or cancern
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/*tag= d
/mod_base= OTHER
/note= "Inverted deoxy abasic nucleotide"
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2002US-0406784P.
2002US-0408378P.
2002US-0409293P.
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2002US-0363124P.
2002US-0386782P.
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29-AUG-2002;
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise riboniclectides and may be double or single stranded. They further comprise sense and antiennse regions, or alternatively are assembled from a sense oligonuclectide and an antisense oligonuclectide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). He siNAs can be unmodified or chemically modified, can contain deoxyribonuclectides, and can be chemically synthesised, can contain a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are
                                         Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiartitic; antiinflammarcory; gene therapy; tablomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERT; DNA-RNA hybrid; ss.
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/note= "Inverted deoxy abasic nucleotide"
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/note= "Inverted deoxy abasic nucleotide"
21. 22
               Human TERC chemically modified siRNA, SEQ ID 554.
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/mod_base= OTHER
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2002US-0409293P.
2003US-0440129P.
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2002US-0386782P.
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modified_base
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05-SEP-2002;
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                                                                                                                                                                                                                 Synthetic.
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Gaps

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148 CCACCGTTCATTCTAGAGC 166

Local Similarity 73.7

Matches

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ADF93827 standard; RNA; 23 BP

26-FEB-2004 (first entry)

ADF93827;

RESULT 260
ADF93827
ID ADF9382
XX
AC ADF9382
XX
DT 26-FEB-

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used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple solerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siMs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents a chemically modified siRNA targeted to the human TERC mRNA
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21. .22
                                                                                                                                                                                                                         Sequence 23 BP; 4 A; 6 C; 2 G; 2 T; 7 U; 2 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human TERC chemically modified siRNA, SEQ ID 546.
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                                                                                                                                                                                                                                                                                                                             146 TICCACCGIICATICIAGA 164
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/mod_base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                              ADF93819 standard; RNA; 23 BP
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11-MAR-2002; 2002US-0363124P
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*tag=
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                                                                                                                                                                                                                                                                        Local Similarity
les 12; Conserva
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modified_base
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                                                                                                                                                                                           transcript.
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                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 261
                                                                                                                                                                                                                                                                                          Matches
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and marisense regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short interfering RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised. Can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and crandulate cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, cued for treating cancer, restenosis, infectious diseases (specifically synthesis and the subject solerosis, lupters and sheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, and solerosis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents a chemically modified siRNA targeted to the human TERC mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                       New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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                                                                                                                                                                                                                                                                                                                                                                               Example 3; SEQ ID NO 546; 145pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           146 TTCCACCGTTCATTCTAGA 164
                   2002US-0396600P.
2002US-0406784P.
2002US-0408378P.
2002US-0409293P.
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                                                              05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409293P.
15-JAN-2003; 2003US-0440129P.
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2002US-0386782P
                                                                                                                                                               (RIBO-) RIBOZYME PHARM INC.
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                                                                                                                                                                                                              Beigelman L;
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Matches 12; Conservative
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                                                                                                                                                                                                              Mcswiggen J,
                                            29-AUG-2002;
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Synthetic

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The invention relates to short interfering nucleic acids (siNh) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonuclectide and antisense oligonuclectide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short interfering RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonuclectides, and can be chemically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modilate expression of the telomerase genes in cells, tissue cused to modilate expression of the telomerase genes in cells, tissue cused to modilate expression of the telomerase genes in cells, tissue cused to modilate expression of the telomerase genes in cells, tissue cused for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus extrhematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents a chemically modified siRNA targeted to the human TERC mRNA, transplants.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
                                                                                                                             "mod_base= Отных
'note= "Pyrimidine bases are 2'-deoxy-2'-fluoro"
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/mod_base= OTHER
/not== "Inverted deoxy abasic nucleotide"
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/mod_base= OTHER
/mode= "Inverted deoxy abasic nucleotide"
21. .22
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                                                                        Location/Qualifiers
                                                                                                                         base≈ OTHER
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mod base= OTHER
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11-MAR-2002; 2002US-0363124P.
06-UUN-2002; 2002US-0386782P.
17-UUL-2002; 2002US-0396600P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0406784P.
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                                                                                                          *tag=
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                                    Homo sapiens
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                     Score 19; DB 1; Length 23;
Pred. No. 1.9e+02;
5; Mismatches 0; Indele
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/mod_base= OTHER
/note= "Inverted deoxy abasic nucleotide"
Sequence 23 BP; 4 A; 7 C; 3 G; 2 T; 5 U; 2 Other;
                                                                                                                                                                                                                               Human TERC chemically modified BiRNA, SEQ ID 548.
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/*tag= c
/mod_base= OTHER
                                                                      148 CCACCGTTCATTCTAGAGC 166
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                                                                                     2 CCACCGUUCAUUCUAGAGC 20
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2002US-0386782P.
2002US-0396600P.
2002US-0406378P.
2002US-0408378P.
                                                                                                                                                         ADF93821 standard; RNA; 23 BP
                      4.2%;
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                       Query Match
Best Local Similarity 73.73
Matches 14; Conservative
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                                                                                                                                                                                                                                                                                                                                                                               Synthetic
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C downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise sense and antisense regions, or alternatively are assembled from a sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and antisense regions, or alternatively are assembled from a sense oligonucleotide and antisense cliquelestranded RNA, micro-RNA (miRNA) and short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised. CC expressed from a vector or enzymatically synthesised. The invention also crelates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue carpanans (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene corpressing a chemically modified siRNA targeted to the human TERC mRNA rransplant
                                                                                                         The invention relates to short interfering nucleic acids (siNA) which
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New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 23 BP; 4 A; 3 C; 7 G; 2 T; 5 U; 2 Other;
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                                                                 Example 3; SEQ ID NO 548; 145pp; English.
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2002US-036782P.

2002US-0406784P.

2002US-0408378P.

2002US-0409378P.

2002US-040129P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 73.7
Matches 14; Conservative
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29-AUG-2002;
05-SEP-2002;
09-SEP-2002;
15-JAN-2003;
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                                                                                                                                                                                                                                                    The invention relates to a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene comprising one or more chemical modifications and each strand of the double-stranded siNA comprises about 21 nucleotides. The siNA of the invention demonstrates antiarteriosclerotic, neuroprotective, notoropic, antiparkinsonian and anticonvulsant activities and may be useful for down-regulating the expression of an endogenous mammalian target gene and therefore in the treatment of any disease or condition that responds to modulation of gene expression or activity in a cell,
                                                                                                                                                                                                                                                                                                                                                                                                         tissue or organism. The disease or condition may include pulmonary diseases such as restenosis, atherosclerosis, Alzheimer's disease, abarkinson's disease, epilepsy, dementia, huntington's disease or amyotrophic lateral sclerosis. Furthermore, the sinA may be utilised for gene therapy applications. The current sequence is that of the sinA DNA-RNA hybrid of the invention.
                                                                                                                                 New double-stranded short interfering nucleic acid molecule, useful for down-regulating the expression of an endogenous mammalian target gene or for treating diseases that respond to modulation of gene expression or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   double-stranded short interfering nucleic acid; siNA; antiarteriosclerotic; neuroprotective; nootropic; antiparkinsonian; anticonvulsant; pulmonary disease; restenosis; atherosclerosis; allatheimer's; Parkinson's; epilepsy; dementia; huntington's; amyotrophic lateral sclerosis; gene therapy; ss; DNA-RNA hybrid; hTR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                     Fosnaugh K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 19; DB 1; Length 23; Pred. No. 1.9e+02;
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                                                   Pavco P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 23 BP; 4 A; 7 C; 3 G; 2 T; 5 U; 2 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   hTR-targeted sinA DNA-RNA hybrid - SEQ ID 606.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                5; Mismatches
                                                                                                                                                                                                                        Example 24; SEQ ID NO 605; 593pp; English.
                                                   Chowrira B,
                                                                      Thompson J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                148 CCACCGTTCATTCTAGAGC 166
               (SIRN-) SIRNA THERAPEUTICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2 CCACCGUUCAUUCUAGAGC 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4.2%;
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2002US-0408378P.
2002US-0409293P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2002US-0363124P.
2002US-0386782P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20-FEB-2003; 2003WO-US005028
                                                     Beigelman L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADG30040 standard; RNA; 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity 73.7
                                                               Usman N,
                                                                                                     WPI; 2003-731676/69.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO2003074654-A2.
                                                     Mcswiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           29-AUG-2002;
05-SEP-2002;
09-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-FEB-2002;
11-MAR-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    26-FEB-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12-SEP-2003
                                                                     Jamison S,
                                                                                                                                                                                      activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADG30040;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 265
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Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAZ07305;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 267
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
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                                                                                                                                                                                                                                                                                                                                                     The invention relates to a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene comprising one or more chemical modifications and each strand of the double-stranded siNA comprises about 21 nuclectides. The siNA of the invention demonstrates antiarteriosclerotic, neuroprotective, nootropic, antipartinsonian and anticonvulsant activities and may be useful for down-regulating the expression of an endogenous mammalian target gene and therefore in the treatment of any disease or condition that responds to modulation of gene expression or activity in a cell, tissue or organism. The disease or condition may include pulmonary diseases such as restenosis, atherosofierosis, Alzheimer's disease. Parkinson's disease, epilepsy, dementia, huntington's disease or congeniem cannot be accounted to a strandary adjacentions. The current sequence is that of the siNA DNA-RNA hybrid of the invention.
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                                                                                                                                                                                                New double-stranded short interfering nucleic acid molecule, useful for down-regulating the expression of an endogenous mammalian target gene or for treating diseases that respond to modulation of gene expression or activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   antiarteriosclerotic; neuroprotective; nootropic; antiparkinsonian; anticonvulsant; pulmonary disease; restenosis; atherosclerosis; Alzheimer's; Parkinson's; epilepsy; dementia; huntington's; amyotrophic lateral sclerosis; gene therapy; ss; DNA-RNA hybrid; hTR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                         Fosnaugh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity 73.7%; Score 19; DB 1; Length 23; Matches 14; Conservative 5; Minmatch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 double-stranded short interfering nucleic acid; siNA;
                                                                                         Pavco P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 23 BP; 4 A; 3 C; 7 G; 2 T; 5 U; 2 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    hTR-targeted sina DNA-RNA hybrid - SEQ ID 604.
                                                                                                                                                                                                                                                                                                                 Example 24; SEQ ID NO 606; 593pp; English.
                                                                                       Chowrira B,
                                                                                       J, Beigelman L, Chowr
, Usman N, Thompson J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     300 GAAGAGTTGGGCTCTGTCA 318
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2 GAAGAGUUGGGCUCUGUCA 20
                                            (SIRN-) SIRNA THERAPEUTICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADG30038 standard; RNA; 23 BP.
15-JAN-2003; 2003US-0440129P.
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06-JUN-2002; 2002US-03867B2P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
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                                                                                                                                                        WPI; 2003-731676/69.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20-FEB-2002;
                                                                                         Mcswiggen J.
Jamison S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12-SEP-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADG30038;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 266
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The invention relates to a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene comprising one or more chemical modifications and each strand of the double-stranded siNA comprises about 21 nucleotides. The siNA of the invention demonstrates antiarteriosclerotic, neuroprotective, nototopic, antiparkinsonian and anticonvulsant activities and may be useful for down-regulating the expression of an endogenous mammalian target gene and therefore in the treatment of any disease or condition target gene and therefore in the treatment of any disease or condition target gene and therefore in the creatment of any disease or condition by the sponds to modulation of gene expression or activity in a cell, tissue or organism. The disease or condition may include pulmonary diseases such as restenosis, atherosclerosis, Alzheimer's disease or Parkinson's disease, epilepsy, dementia, huntington's disease or emperatoric planteral sclerosis. Furthermore, the siNA may be utilised for gene therapy applications. The current sequence is that of the siNA DNA-RNA hybrid of the invention.
                                                                                                                                                                                                                                                                                                                                                                   New double-stranded short interfering nucleic acid molecule, useful for down-regulating the expression of an endogenous mammalian target gene or for treating diseases that respond to modulation of gene expression or activity.
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                                                                                                                                                                                     Fosnaugh K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human telomerase RNA gene (hTR) promoter specific primer hlle.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4.2%; Score 19; DB 1; Length 23;
53.2%; Pred. No. 1.9e+02;
Ive 7; Mismatches 0; Indels
                                                                                                                                                                                 Pavco P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 23 BP; 4 A; 6 C; 2 G; 2 T; 7 U; 2 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 24; SEQ ID NO 604; 593pp; English.
                                                                                                                                                                                 Chowrira B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RES CAMPAIGN TECHNOLOGY.
                                                                                                                                                                                                                        Thompson J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     146 TICCACCGTICATICIAGA 164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ::||||||::||::|:||
uuccacceuucauucuaga 20
                                                                                                            (SIRN-) SIRNA THERAPEUTICS INC.
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09-SEP-2002; 2002US-0409293P.
15-JAN-2003; 2003US-0440129P.
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                                                                                                                                                                                     Beigelman L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAZ07305 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22-OCT-1999 (first entry)
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nes 12, Conservative
                                                                                                                                                                                                                        Usman N,
                                                                                                                                                                                                                                                                                             WPI; 2003-731676/69
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                                                                                                                                                                                     Mcswiggen J,
Jamison S, t
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Page 140

US09436060A-16.rng.sl

Disclosure, Fig 6; 109pp; English.

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The invention relates to promoter regions from mouse and human telomerase

RNA (TR) component genes. The TR gene promoter can be linked to a

Reterologous gene, especially a gene encoding a cytocoxin, for therapy of

cancer, especially neoplasias. The telomerase is necessary for the

unrestricted proliferative capacity of many human cancers. Mutation or

cyrregulation of the telomerase expression pathway may cause reactivation

or upregulation of telomerase expression in cancer. Substances,

identified in the methods, can be used to block transcription from the TR

gene promoter through interaction of the 5 regulatory sequences. These

substances, e.g. antisense oligonocleotides, transcription factors,

peptide nucleic acids and factors that disrupt signal transduction, are

useful for cancer therapy. In particular, gene therapy vectors sequences

(specially pGT62-codAupp) comprising the promoter and a viral thymidine

kinase gene can be controlled or treated. Direct down-regulation of

telomerase RNA gene through manipulation of transcription factors may be

effective anticancer therapy and the cloning of the hTR gene promoter

callows the analysis of therappeutic molecules which modulate hTR promoter

activity, Sequences AAZ07656-321 regules

and mutagenisis of human TR gene (HTR) promoter region
                                     Mouse and human telomerase RNA gene promoters, useful for tumor specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Telomerase RNA; TR; promoter; cytotoxin; cancer; neoplasia; terc; gene therapy; thymidine kinase gene; anticancer therapy; mouse;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 22 BP; 3 A; 4 C; 12 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mouse telomerase RNA gene specific primer mTRr1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Mismatches
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                                                                                       Disclosure; Fig 12; 109pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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Les 20; Conservat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1999-479183/40.
WPI; 1999-479183/40
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                                                       gene therapy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Keith WN;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 268
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Mouse and human telomerase RNA gene promoters, useful for tumor specific

gene therapy

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Gaps

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The invention relates to promoter regions from mouse and human telomerase CC RNA (TR) component genes. The TR gene promoter can be linked to a heterologous gene, especially a gene encoding a cytotoxin, for therapy of cancer, especially neoplasias. The telomerase is necessary for the curestricted proliferative capacity of many human cancers. Mutation or unrestricted proliferative expression pathway may cause reactivation of cypregulation of telomerase expression in cancer. Substances, cor upregulation of telomerase expression in cancer. Substances, cor upregulation of telomerase expression in cancers. Substances, cor upregulation of telomerase expression of the 5' regulatory sequences. These substances, e.g. antisense oligonucleotides, transcription factors, cypetide nucleic acids and factors that disrupt signal transduction, are useful for cancer therapy. In particular, gene therapy vectors (especially pGT62-codAupp) comprising the promoter and a viral thymidine converted to convert a profucy. Gene promoter and a viral thymidine conpasia can be controlled or transcription factors may be effective anticancer therapy and the cloning of the hrm gene promoter callows the analysis of therapeutic molecules which modulate hrm promoter callows the analysis of therapeutic molecules which modulate hrm promoter cativity. Sequences AAZ07681-95 represent PCR primers for amplifying consecurations.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to promoter regions from mouse and human telomerase RNA (TR) component genes. The TR gene promoter can be linked to a heterologous gene, especially a gene encoding a cytotoxin, for therapy of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mouse and human telomerase RNA gene promoters, useful for tumor specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Telomerase RNA; TR; promoter; cytotoxin; cancer; neoplasia; terc; gene therapy; thymidine kinase gene; anticancer therapy; mouse;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4.1%; Score 18.4; DB 1; Length 20; 95.0%; Pred, No. 1.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 6 A; 5 C; 7 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mouse telomerase RNA gene specific primer mTRrl.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (CANC-) CANCER RES CAMPAIGN TECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Fig 6; 109pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        102 TTCTCGCTGACTTTCAGCGG 121
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Best Local Similarity 95.0°
Matches 19; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  qene therapy.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAZ07294;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Keith WN;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 269
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unrestricted proliferative capacity of many human cancers. Mutation or dysregulation of the telomerase repression pathway may cause reactivation of repression pathway may cause reactivation or upregulation of telomerase expression pathway may cause reactivation or upregulation of telomerase expression in cancer. Substances, dentified in the methods, can be used to block transcription from the TR gene promoter through interaction of the 5' regulatory sequences. These substances, e.g. antisense oligonucleotides, transcription factors, peptide nucleic acids and factors that disrupt signal transduction, are useful for cancer therapy. In particular, gene therapy vectors (especially prof62-codhupp) comprising the promoter and a viral thymidine kinase gene can be used to convert a prodrug, e.g. gancyclovir, so that neoplasia can be used to convert a prodrug, e.g. gancyclovir, so that coplasia can be used to convert a prodrug, e.g. gancyclovir, so that neoplasia can be through manipulation of transcription factors may be effective anticancer therapy and the cloning of the hTR gene promoter allows the analysis of therapeutic molecules which modulate hTR promoter activity. Sequences AAZ07681-95 represent PCR primers for amplifying
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Telomerase; human; cancer; diagnosis; melanoma; skin cancer; leukaemia; neuroblastoma; breast carcinoma; colon carcinoma; lymphoma; osteosarcoma; smooth muscle cell hyperplasia; stem cell proliferation; Wilm's tumour; stem cell differentiation; organ differentiation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    products for
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4.1%; Score 18.4; DB 1; Length 20; 95.0%; Pred. No. 1.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New isolated vertebrate telomerase genes - used to develop
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Seguence 20 BP; 6 A; 5 C; 7 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   gene (terc) promoter sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      102 TTCTCGCTGACTTTCAGCGG 121
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97US-0053018P.
97US-0053329P.
97US-0054642P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PCR primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
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This sequence is a PCR primer for DNA encoding a truncated human

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telomerase of the invention. Primers that amplify the telomerase coding sequence can be used in a method for diagnosing cancer in a patient. The telomerase can be used for detection, diagnosis and drug screening. Inhibitors of telomerase activity can be used to treat cancers such as melanomas, other skin cancers, neuroblastomas, breast carcinomas, colon carcinomas, leukaemias, lymphomas, osteosarcomas or smooth muscle cell hyperplasias or skin growths. Enhancers of telomerase may be used to stimulate stem cells proliferation and differentiation (expansion of haematopoietic stem cells could be administered in the bone marrow transplant context). As well, many tissues have stem cells. Proliferation of these cells may be useful in wound healing, hair growth, treatment of disease such as Wilm's tumour, organ regeneration or differentiation after injury or diseases, nerve cell or brain cell growth following
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New peptide nucleic acid (PNA) compounds that inhibit telomerase activity in mammalian cells is useful as probes to detect the RNA component of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note = "Peptide nucleic acid molecule, where N-(2-aminoethyl)glycine units are linked to nucleotide bases via glycine amino N through a methylenecarbonyl linker"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present sequence represents a peptide nucleic acid molecule which hybridises to the mRNA component of mammalian telomerase, and inhibits telomerase activity. Telomerase is a ribonucleoprotein enzyme that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Peptide nucleic acid; PNA; telomerase; ribonucleoprotein enzyme; cance inhibitor; neoplasia; neurodegenerative diesase; aging; hyperplasia; AIDS; HIV; fungal infection; forensic identification; detect; tumour; pacernicy testing; ss.
                                                                                                                                                                                                                                                                                                                                      Gaps
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                                                                                                                                                                                                                                                                                                  Score 18; DB 1; Length 18;
Pred. No. 1.7e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PNA sequence #9 used to inhibit telomerase activity.
                                                                                                                                                                                                                                                                   Sequence 18 BP; 1 A; 2 C; 12 G; 3 T; 0 U; 0 Other;
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100.0%; Pred. No. ...
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                                                                                                                                                                                                                                                                                                                                  18; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     mammalian telomerase.
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                                                                                                                                                                                                                                                                                                                      Similarity
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Best Local 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 271
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synthesizes one strand of the telomeric DNA, using as a template an 11 nucleotide sequence contained within the RNA component of the enzyme. The invention relates to PNA molecules having a sequence of no more than 25 bases, which include the sequence of TTAGG. The uncharged nature of the PNA carecombinates the melting temperature of associating strands, increases the rate of association with targeted nucleic acids, and affords greater resistance of degradation by proteases or nucleases. The therapeutic PNAs may be used for treating disease conditions such as cancers, neoplasia, hyperplasia, neurodegenerative diseases, aging, human immunodeficiency virus (HIV) infection/AIDS (acquired immunodeficiency virus activity, in combination with antineoplastic and other cytomerase RNA (HIR) sequence. PNA probes are also used for forensic identification of individuals, e.g. paternity testing, based on hTR gene restriction of individuals, e.g. paternity testing, based on hTR gene restriction individuals, e.g. paternity testing, based on hTR gene actriction and hold of the present invention allows cancerous conditions of a annewous conditions to be detected with increase and as annewous conditions are also used for the present invention allows the page and as allows cancerous conditions to be detected with increases and as annewous enditions are accounted to a manner and the present invention and and the p
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//note= "This sequence is a peptide nucleic acid, i.e. it
contains a polyamide backbone instead of a deoxyribose
backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           possibly at an earlier stage, before cells are detected as cancerous based on pathological characteristics. The diagnostic and prognostic methods of the present invention can be used to detect an immortal or neoplastic cell or tumour tissue or cancer of any origin, provided the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mammalian; peptide nucleic acid; probe; forensic; paternity testing; human telomerase RNA component; hTR gene RFLP pattern; cancer; inflammation; lymphoproliferative disease; autoimmune disease; neurodegenerative disease; neoplasia; hyperplasia; HIV; AIDS; telomera metabolism; mutant; acquired immunodeficiency syndrome; telomere metabolism; mutant; cytostatic; anti-inflammatory; immunosuppressive; polyamide backbone; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PNA 27 inhibiting human and mammalian telomerase activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 18; DB 1; Length 18;
Pred. No. 1.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                expresses telomerase activity and its RNA component
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 18 BP; 2 A; 5 C; 4 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Match
Local Similarity 100.0%; Pr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AACCCTAACTGAGAAGGG 65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18 AACCCTAACTGAGAAGGG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAS15430 standard; DNA; 18 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Key
modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           14-FEB-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 272
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The present invention relates to peptide nucleic acids (PNAs), comprising a sequence of 6-25 nucleobases, that inhibit telomerase activity in mammalian cells by hybridising to the RNA component of mammalian cells by hybridising to the RNA component of mammalian cells by hybridising to the RNA component of mammalian telomerase and as inhibitors of telomerase activity, or to detect and/or quantitate polynucleotide having the human telomerase RNA component (hTR) sequence, as well as in forensic identification of individuals, such as paternity testing or identification of cindividuals, such as paternity testing or identification of criminal suspects or unknown descendants based on the hTR gene RFLP pattern. The Suppects or unknown descendants based on the hTR gene RFLP pattern. The CTP MA can be further used for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases. The PNAs in combination with other pharmaceuticals (such as antineoplastic or cytostatic agents) can be used for treating neoplasia, hyperplasia, human immunodeficiency virus (HTV) infections, acquired immunodeficiency syndrome (AIDS) and associated pathologies, and other diseases characterised by abnormal telomere metabolism or telomerase of the invantion.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                 New peptide nucleic acids that hybridizes to the RNA component of mammalian telomerase, useful for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Telomerase RNA; TR; promoter; cytotoxin; cancer; neoplasia; hTR; gene therapy; thymidine kinase gene; anticancer therapy; human; mutagenesis; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human telomerase RNA gene (hTR) promoter specific primer h112b.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 18 BP; 2 A; 5 C; 4 G; 7 T; 0 U; 0 Other;
                                                                                             Wright WE, Piatyszek MA, Corey DR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (CANC-) CANCER RES CAMPAIGN TECHNOLOGY.
                                                                                                                                                                                                                                                                         Claim 7; Col 73; 46pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              48 AACCCTAACTGAGAAGGG 65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18 AACCCTAACTGAGAAGGG 1
                97US-00838545.
96US-00630019.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22-OCT-1999 (first entry)
                                                    (TEXA ) UNIV TEXAS SYSTEM.
                                                                                                                                 WPI; 2001-638024/73
09-APR-1996;
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                09-APR-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      activity. 7
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                                                                                               Shay JW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
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Matches
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ROBB SA;
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                                                                                                                                      The invention relates to promoter regions from mouse and human telomerase
RNA (TR) component genes. The TR gene promoter can be linked to a
cheterologous gene, sepecially a gene encoding a cytocoxin, for therapy of
cheterologous gene, sepecially a gene encoding a cytocoxin, for therapy of
corner, especially nooplasias. The telomerase is necessary for the
unrestricted proliferative capacity of many human cancers. Mutation or
dysregulation of the telomerase expression pathway may cause reactivation
c or upregulation of telomerase expression pathway may cause reactivation
c or upregulation of telomerase expression pathway may cause reactivation
c upregulation of telomerase expression pathway may cause reactivation
c especially no cancer through interaction of the 5' regulatory sequences. These
cuseful for cancer therapy. In particular, gene therapy vectors
c especially pGT62-codAupp) comprising the promoter and a viral thymidine
c especially pGT62-codAupp) comprising the promoter and a viral thymidine
c kinase gene can be used to convert a prodrug, e.g. gancyclovir, so that
neoplasia can be used to convert a prodrug, e.g. gancyclovir, so that
neoplasia can be used through manipulation of transcription features may be
telomerase RNA gene through manipulation of transcription features may be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
                                                              Mouse and human telomerase RNA gene promoters, useful for tumor specific
                                                                                                                                                                                                                                                                                                                                                                                                            effective anticancer therapy and the cloning of the hTR gene promoter allows the analysis of therapeutic molecules which modulate hTR promoter activity. Sequences AAZ07696-321 represent PCR primers used in cloning and mutagenisis of human TR gene (hTR) promoter region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RNA component; human telomerase; antisense oligonucleotide; infection; neuroblastoma; bladder cancer; colon cancer; prostate cancer; cancer; contraception; sterilisation; immunosuppression; therapeutic; hTR; immune system down-regulation; anti-inflammatory therapy; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RNA component of human telomerase (hTR) antisense oligo RP2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3.8%; Score 17.2; DB 1; Length 22;
86.4%; Pred. No. 2.5e+02;
ive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pruzan R, Weinrich SL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 22 BP; 2 A; 4 C; 12 G; 4 T; 0 U; 0 Other;
                                                                                                                Disclosure; Fig 12; 109pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15 GGGCCTGGGAGGGGTGGTGGCC 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 ececciceeraacinericeic 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAV41181 standard; DNA; 17 BP.
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96US-00770565
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
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Best Local Similarity 86.4
                                WPI; 1999-479183/40.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (GERO-) GERON CORP.
                                                                                 gene therapy.
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20-DEC-1996;
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 Keith WN;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 274
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Sequences shown in AAV41169 to AAV41181 represent antisense
oligonucleotides to the RNA component of human telomerase (hTR). These
antisense oligonucleotides specifically hybridise to a nucleotide
caquence within an accessible region of the hTR, but that does not
hybridise to a sequence within the template region of hTR. These
cligonucleotides may specifically be used for detection of an RNA
component of human telomerase in a sample. This is useful for diagnosing
cancer (especially neuroblastoma, bladder, colon and prostate cancer),
and providing prognosis for a cancer patient. The inhibitory
clife of the telomerase RNA component, decreasing the half-
interfering with transcription of the RNA component, decreasing the half-
clife of the telomerase RNA component transcript, inhibiting assembly of
the RNA component into the telomerase notivery in both cultured cells and in
be used for inhibiting telomerase activity in both cultured cells and in
be used for inhibiting telomerase activity in both cultured cells and in
clease inhibiting telomerase activity in both cultured cells and in
preventing cancer, for contraception or sterilisation, for
immunosuppression, and for selectively down-regulating specific branches
cof the immune system, e.g. a specific subset of T-cells, in anti-
inflammatory therapies or for treating infections by, e.g. yeast,
parasites or fungi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
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  - used for detecting
New polynucleotide(s) anti:sense to human telomerase - used for detecting or inhibiting human telomerase, e.g. for treating cancers, contraception, immuno-suppression or treating infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         acyl-coenzyme A synthetase 1; ACS1; diabetes; obesity; metabolic syndrome X; cardiovascular disorder; cancer; infection; inflammation; tumour; antisense; 88.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3.8%; Score 17; DB 1; Length 17; 100.0%; Pred. No. 1.9e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; Prec. ...
                                                                                                                  Claim 11; Page 65; 80pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       177 TGTCAGCTGCTGGCCCG 193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP.
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Matches 17; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2004-203782/19.
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Sequence 20
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                                                                                                                                                                                                                                                            ADK20636;
                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ross SA;
                                                                                                       Best Loca
Matches
                                                                                                                                                                                                         RESULT 277
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                                                                                                                                                                                                                                                                           ö
                                               The invention relates to an antisense compound targeted to a nucleic acid molecule encoding acyl-coenzyme A synthetase 1 (ACS1). The antisense compound specifically hybridises with and inhibits the expression of ACS1. The antisense oligonucleotides or compounds are useful for inhibiting the expression of acyl-coenzyme A synthetase 1 (ACS1), and for treating diseases or conditions associated with aberrant expression of ACS1, e.g. diabetes, obesity, metabolic syndrome X, cardiovascular disorder or cancer. The antisense compounds are also useful as research reagents and kits, or in diagnostic, therapeutic and prophylactic applications, e.g. to prevent or delay infection, inflammation or tumour formation. The present sequence represents an acyl-coenzyme A synthetase 1, ACS1, antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to an antisense compound targeted to a nucleic acid molecule encoding acyl-coenzyme A synthetase 1 (ACS1). The antisense compound specifically hybridises with and inhibits the expression of ACS1. The antisense oligonuclectides or compounds are useful for inhibiting the expression of acyl-coenzyme A synthetase 1 (ACS1), and for treating diseases or conditions associated with aberrant expression of ACS1, e.g. diabetes, obesity, metabolic syndrome X, cardiovascular disorder or cancer. The antisense compounds are also useful as research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New antisense compounds targeted to nucleic acid molecules encoding acylcoenzyme A synthetase 1 (ACS1), useful for treating diseases or conditions associated with aberrant expression of ACS1, e.g. diabetes,
                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Acyl-coenzyme A synthetase 1, ACS1, antisense oligonucleotide #566.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 acyl-coenzyme A synthetase 1; ACS1; diabetes; obesity; metabolic syndrome X; cardiovascular disorder; cancer; infection; inflammation; tumour; antisense; ss.
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                                                                                                                                                                                                                                           Score 16.4; DB 1; Length 20;
Pred. No. 2.6e+02;
0; Mismatches 1; Indels
                                                                                                                                                                                                                       Sequence 20 BP; 1 A; 12 C; 4 G; 3 T; 0 U; 0 Other;
                         Claim 3; SEQ ID NO 632; 940pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 3; SEQ ID NO 566; 940pp; English.
                                                                                                                                                                                                                                                                                                   349
                                                                                                                                                                                                                                                                                                                                                                                             BP
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                                                                                                                                                                                                                                                 3.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  14-AUG-2003; 2003WO-US025389
                                                                                                                                                                                                                                                                                                    CGGGGGCGAGGGCGAGGT
                                                                                                                                                                                                                                                                                                                             CGAGGGCGAGGCGAGGT
                                                                                                                                                                                                                                                                                                                                                                                             ADK20489 standard; DNA; 20
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                                                                                                                                                                                                                                                                         17; Conservative
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                                                                                                                                                                                                                                                              Local Similarity
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 obesity or cancer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                                                                                    332
                                                                                                                                                                                                                                                                                                                             13
                                                                                                                                                                                                                                                                                                                                                                                                                       ADK20489;
                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ross SA;
                                                                                                                                                                                                                                                                           Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to an antisense compound targeted to a nucleic acid molecule encoding acyl-coenzyme A synthetase 1 (ACSI). The antisense compound specifically hybridises with and inhibits the expression of ACSI. The antisense oligonuclectides or compounds are useful for inhibiting the expression of acyl-coenzyme A synthetase 1 (ACSI), and for treating diseases or conditions associated with aberrant expression of ACSI, e.g. diabetes, obesity, metabolic syndrome X, cardiovascular disorder or cancer. The antisense compounds are also useful as research reagents and kite, or in diagnostic, therapeutic and prophylactic applications, e.g. to prevent or delay infection, inflammation or tumour formation. The present sequence represents an acyl-coenzyme A synthetase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New antisense compounds targeted to nucleic acid molecules encoding acylocarzyme A synthetase 1 (ACS1), useful for treating diseases or conditions associated with aberrant expression of ACS1, e.g. diabetes,
reagents and kits, or in diagnostic, therapeutic and prophylactic applications, eg. to prevent or delay infection, inflammation or tumour formation. The present sequence represents an acyl-coenzyme A synthetase 1, ACS1, antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Acyl-coenzyme A synthetase 1, ACS1, antisense oligonucleotide #713.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     acyl-coenzyme A synthetase 1; ACS1; diabetes; obesity;
metabolic syndrome X; cardiovascular disorder; cancer; infection;
inflammation; tumour; antisense; ss.
                                                                                                                                                                                                                                                                                                                           0;
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0
                                                                                                                                                                                                                                                3.6%; Score 16.4; DB 1; Length 20;
94.4%; Pred. No. 2.6e+02;
ve 0; Mismatches 1; Indels
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14.4%; Pred. No. 2.6e+02;
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                                                                                                                                                                                BP; 1 A; 12 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        332 CGGGGGGGGGGGGGT 349
                                                                                                                                                                                                                                                                                                                                                                                                      332 CGGGGGCGAGGCGAGGT 349
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18 ccacccacccacccacct 1
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                                                                                                                                                                                                                                                                                  Local Similarity 94.4%;
nes 17; Conservative
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Best Local Similarity 94.4%;
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADK20636 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
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The inventors describe the production of mammalian cells expressing prohormone convertase which facilitates the processing of prohormone produces to active hormones. More specifically the cleavage site is the prohormone convertase cleavage site given in ARR37632-35. In order to define murine prohormone convertase 1 specifically prorelaxin alanine mutations of the basic residues K and R were constructed. The following mutations were tested for substrate specificity to murine prohomone convertase: 1.4, 2.4, 3.2, 4.3, 7.2 and 8.6 (see AAQ43256-63; AAR37619-26). (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence encoding the C-chain/A-chain junction prorelaxin (Rx) mutant Rx 2.4 at the prohormone convertage cleavage site.
                                                                                              Sequence encoding the C-chain/A-chain junction prorelaxin (Rx) mutant Rx 1.4 at the prohormone convertase cleavage site.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Heterologous polypeptide factor prepn. - by introducing into polypeptide factor dependent host cell nucleic acid, and then culturing host cell,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Prohormone convertage; enzyme; cleavage; prorelaxin; hormone;
                                                                                                                                                     Prohormone convertase; enzyme; cleavage; prorelaxin; hormone;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
3.6%; Score 16.2; DB 1; Length 21;
Best Local Similarity 85.7%; Pred. No. 2.8e+02;
Matches 18; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 21 BP; 10 A; 4 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Marriott D;
                                                                                                                                                                                                                                     Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       156 CATTCTAGAGCAAACAAAAA 176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 CATTCTAGAGCAAAGAGACAA 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example; Page 54; 134pp; English.
                                                                                                                                                                                                                                                                                                                                                                                             92WO-US010621.
                                                                                                                                                                                                                                                                                                                                                                                                                                  91US-00803631.
92US-00887265.
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                                      (revised)
(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (GETH ) GENENTECH INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1993-197065/24.
P-PSDB; AAR37620.
                                                                                                                                                                                                                                                                                                               WO9311247-A1
                                                                                                                                                                                                                                                                                                                                                                                             04-DEC-1992;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      22-MAY-1992;
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24-SEP-1993
                                      25-MAR-2003
24-SEP-1993
                                                                                                                                                                                                                                                                                                                                                      10-JUN-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gorman CM,
                                                                                                                                                                                                Synthetic.
AAQ43257;
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                                                                                                                                                                                                                                     Key
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  XXXXEEXAXXXXX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2 Primers (see AAV27886 and AAV27887) are used in the PCR amplification of a 520 bp fragment of telomerase genomic DNA from HeLa cells. The PCR product was used to prepare 2 DNA constructs, each containing the T7 promoter. One construct (for primers see AV27888-89) contained DNA which would generate sense strand human telomerase RNA in a transcription reaction. The other construct (for primers see AAV27890-91) contained DNA which would generate antisense strand human telomerase RNA in a transcription reaction. The human telomerase RNA in a transcription reaction. The human telomerase RNA was used for in vitro assays to demonstrate that wild-type TP2 and sense telomerase RNA are required for telomerase activity, Novel TP2 can be used to develop products that regulate telomerase activity, useful e.g. for treating fancer, AIDS and ageing disorders. (Updated on 25-MAR-2003 to correct PR field.)
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                                                                                                                                                                                                                                                                                            PP2; human; telomerase protein 2; cancer; AIDS; ageing; therapy; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New nucleic acid encoding human telomerase protein-2 - used for regulating telomerase activity, e.g. for treating cancer or acquired
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 21 BP; 1 A; 5 C; 13 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 8; Page 90; 150pp; English.
                                                                                                                                                                                                                                                          Human telomerase gene PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 GGGTTGCGGAGGGTGGGC 18
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                                                                                                                    AAV27886 standard; DNA; 21 BP
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97US-00951733,
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Best Local Similarity 94.4%;
Marches 17; Conservative
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CGAGGGCGAGGCGAGGT
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                                                                                                                                                                                                (revised)
(first entry)
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(AMGE-) AMGEN CANADA INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1998-297946/26.
                                                                                                                                                                                                                                                                                                                                                      Synthetic.
Homo Bapiens.
                                                                                                                                                                                                                                                                                                                                                                                                               WO9821343-A1
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16-OCT-1997;
                                                                                                                                                                                              25-MAR-2003
12-OCT-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               13-NOV-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        22-MAY-1998
                                                                                                                                                                                                                                                                                                                     primer; ss
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RESULT 279 AAQ43257 ID AAQ432 XX

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A series of natural or mutated cDNAs for the prohormone convertase (PC) cleavage sites in human proralaxin protein between the C and A chains (AAQ71455-59 corresponding to AAR60583-7) or in the B and C chains (AAQ71460-63 corresponding to AAR60588-90). The natural cleavage sites contain two dibasic residues, Lys-Arg, both of which are required for cleavage. The mutant sites have replacements of either residues with an Ala. This confers specificity to murine PCI. In this particular mutant, the Lys residue at the fourth position of the cleavage site has been responsible for remodelling the reproductive tract prior to parturition. The cells containing both the prohormone convertase gene and the required precursor to be expressed can be injected into a mammal. (Updated on 25-
                                                                                                                                Treating insulin-dependent disorders in mammals - by introducing a nucleic acid encoding a variant proinsulin into a host cell with a constitutive pathway of protein secretion, or a plasmid, and introducing the cell or plasmid to the mammal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       neuroblastoma; breast carcinoma; colon carcinoma; lymphoma; osteosarcoma; smooth muscle cell hyperplasia; stem cell proliferation; Wilm's tumour; stem cell differentiation; organ regeneration; organ differentiation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Telomerase; human; cancer; diagnosis; melanoma; skin cancer; leukaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 3.6%; Score 16.2; DB 1; Length 21; Best Local Similarity 85.7%; Pred. No. 2.8e+02; Matches 18; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 21 BP; 10 A; 4 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PCR primer for telomerase coding sequence.
                                                                                                                                                                                                                      Example 3; Page 52; 117pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   156 CATTCTAGAGCAAACAAAAA 176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 carrcragagcaaagagaga 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MAR-2003 to correct PN field.)
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97US-0053018P.
97US-0053329P.
97US-0054642P.
97US-0058287P.
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                                            Groskreutz DJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAX18326 standard; DNA; 21
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     (GETH ) GENENTECH INC.
                                                                        WPI; 1994-303031/37.
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                                                                                              P-PSDB; AAR60585
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PCR primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               26-JUL-1999
                                        Gorman CM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-JUL-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-JUL-1997,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21-JUL-1997;
21-JUL-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           04-AUG-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kilian A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAX18326;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The inventors describe the production of mammalian cells expressing prohormone convertase which facilitates the processing of prohormone products to active hormones. More specifically the cleavage site is the prohormone convertase cleavage site given in ARR3762-35. In order to define murine prohormone convertase 1 specifically prorelaxin alanine mutations of the basic residues K and R were constructed. The following mutations were tested for substrate specificity to murine prohomone convertase: 1.4, 2.4, 3.2, 4.3, 7.2 and 8.6 (see AAQ43256-63; AAR37619-26). (Updated on 25-WAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                          Heterologous polypeptide factor prepn. - by introducing into polypeptide factor dependent host cell nucleic acid, and then culturing host cell,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           prohormone; convertase; insulin; proinsulin; factor; growth; precursors;
transgenic; mammal; prorelaxin; cleavage; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3.6%; Score 16.2; DB 1; Length 21;
85.7%; Pred. No. 2.8e+02;
ive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Rx 2.4 prorelaxin C/A chain junction mutant cleavage site.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 21 BP; 10 A; 4 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                      Gorman CM, Groskreutz DJ, Marriott D;
                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CATTCTAGAGCAAACAAAAA 176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CATTCTAGAGCAAAGAGACAA 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example; Page 54; 134pp; English.
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                                                                                                                                                                               92WO-US010621
                                                                                                                                                                                                                91US-00803631
                                                                                                                                                                                                                                   92US-00887265
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(first entry)
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                                                                    /*tag=
                                                                                                                                                                                                                                                                  (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                         WPI; 1993-197065/24.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
                                                                                                                                                                                                                                                                                                                                                          P-PSDB; AAR37621
                                                                                                        WO9311247-A1
                                                                                                                                                                             04-DEC-1992;
                                                                                                                                                                                                            06-DEC-1991;
                                                                                                                                                                                                                                 22-MAY-1992;
                                                                                                                                         10-JUN-1993
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-MAR-1993;
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26-APR-1995
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Synthetic
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                                                                                                                                        This sequence is a PCR primer for DNA encoding a truncated human telenomerase of the invention. Primers that amplify the telenomerase coding sequence can be used in a method for disquesing cancer in a patient. The telenomerase can be used for detection, disquesis and drug screening. Inhibitors of telenomerase activity can be used to treat cancers such as meananomas, other skin cancers, neuroblastomas, breast cancers such as carcinomas, lowthere skin cancers, neuroblastomas, breast cancers such as hyperplasias or skin growths. Enhancers of telemerase may be used to stimulate stem cells proliferation and differentiation (expansion of haematopoietic stem cells could be administered in the bone marrow transplant context). As well, many tissues have stem cells. Proliferation of these cells may be useful in wound healing, hair growth, treatment of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              used
                                             New isolated vertebrate telomerase genes - used to develop products for treating cancers or for organ regeneration, nerve cell or brain cell growth following injury or bone marrow transplantation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                       disease such as Wilm's tumour, organ regeneration or differentiation after injury or diseases, nerve cell or brain cell growth following
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Peptide nucleic acid; PNA; cancer; telomerase; probe; hybridisation;
inhibitor; human telomerase RNA; hTR; PCR; oligonucleotide; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            , peptide nucleic acids hybridising to mammalian telomerase RNA - inhibit telomerase, for treating tumours and other proliferative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA oligonucleotide 3, used in the measurement of Tm values.
                                                                                                                                                                                                                                                                                                                                                                                                                                      3.6%; Score 16.2; DB 1; Length 21; 85.7%; Pred. No. 2.8e+02; ive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Norton JC;
                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 21 BP; 3 A; 5 C; 9 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ď,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Corey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Piatyszek MA,
                                                                                                             Example 1; Page 43; 134pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       431 CAGGACTCGCCTCACACATGC 451
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 2; Page 49; 76pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CAGGACTCGGCTCACCACTGC 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              diseases, also for diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAT89247 standard; DNA; 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Shay JW, Wright WE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1997-512647/47.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (GERO-) GERON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21
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                                                                                                                                                                                                                                                                                                                                                                            injury
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This is an oligonucleotide used in the measurement of Tm values and their complimentary peptide nucleic acids (PNAs), (e.g. AAT89236-T89239). PNAs

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New peptide nucleic acid (PNA) compounds that inhibit telomerase activity in mammalian cells is useful as probes to detect the RNA component of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Peptide nucleic acid; PNA; telomerase; ribonucleoprotein enzyme; cancer; inhibitor; neoplasia; neurodegenerative disease; aging; hyperplasia; AIDS; HIV; fungal infection; forensic identification; detect; tumour; paternity testing; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note= "Peptide nucleic acid molecule, where N-(2-aminoethyl)glycine units are linked to nucleotide bases via glycine amino N through a methylenecarbonyl linker"
hybridise specifically to an RNA component of mammalian telomerase, and include the sequence GGG for specific hybridisation to the template region of this component. PNAs can be used as probes to detect the RNA component of mammalian telomerase and as inhibitors of telomerase activity, especially in the treatment of cancer
                                                                                                                                                                                             Gaps
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                                                                                                                                                     DB 1; Length 16; 2.1e+02;
                                                                                                                                                                                             0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PNA sequence #27 used to inhibit telomerase activity.
                                                                                                                   Sequence 16 BP; 5 A; 2 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Norton JC,
                                                                                                                                                     3.5%; Score 16; DB
100.0%; Pred. No. 2.1
rative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                    BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       97US-00838545.
                                                                                                                                                                                                                                                              TAACTGAGAAGGGCGT 16
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                                                                                                                                                                                                                                  68
                                                                                                                                                                                                                                                                                                                                                                    AAA37569 standard; DNA; 16
                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
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                                                                                                                                                                                                                                  53 TAACTGAGAAGGGCGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (TEXA ) UNIV TEXAS SYSTEM
                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      mammalian telomerase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2000-292432/25.
                                                                                                                                                       Query Match
Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       09-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              misc feature
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                          AAA37569;
                                                                                                                                                                                                                                                                                                                                RESULT 284
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immunodeficiency virus (HIV) infection/AIDS (acquired immunodeficiency Mammalian; paternity testing; human telomerase RNA component; 3.5%; Score 16; DB 1; Length 16; 100.0%; Pred. No. 2.1e+02; live 0; Mismatches 0; Indels Norton JC; Sequence 16 BP; 5 A; 2 C; 6 G; 3 T; 0 U; 0 Other; Piatyszek MA, Corey DR, AAS15447 standard; DNA; 16 BP 68 1 TAACTGAGAAGGGCGT 16 99US-00349532. 97US-00838545. 96US-00630019, (first entry) 53 TAACTGAGAAGGGCGT (TEXA) UNIV TEXAS SYSTEM. Local Similarity 100. 18s 16; Conservative Shay JW, Wright WE, WPI; 2001-638024/73. US6294650-B1 14-FEB-2002 sapiens 08-JUL-1999; 09-APR-1996; 39-APR-1997; 25-SEP-2001, Synthetic. AAS15447; Query Match RESULT 285 Best Loc Matches Ношо AAS15447 \$ 8 셤

syndrome) and associated by abnormal telomere metabolism or telomerase diseases characterized by abnormal telomere metabolism or telomerase activity, in combination with antineoplastic and other cytooxic or cytostatic agents, antifungal agents, and other nucleotides. PNAS may be cytostatic agents, antifungal agents, and other nucleotides. PNAS may be used for molecular diagnostics, labelled PNAS are used as hybridization probes to detect or quantitate polymucleotides having a human telomerase or a molecular diagnostics, labelled PNAS are used as hybridization probes to detect or quantitate polymucleotides having a human telomerase of individuals, e.g. paternity testing, based on hTR gene restriction of individuals, e.g. paternity testing, based on hTR gene restriction fragment length polymorphism (RFLP) pattern. PNAS are also useful as probes to detect the RNA component of a mammalian telomerase and as inhibitors of telomerase activity. The method of the present invention possibly at an earlier stage, before cells are detected as cancerous conditions to be detected and prognostic based on pathological characteristics. The diagnostic and prognostic methods of the present invention can be used to detect an immortal or neoplastic cell or tumour tissue or cancer of any origin, provided the expresses telomerase activity and its RNA component. ; 0 Gaps hTR gene RFLP pattern; cancer; inflammation; forensic; lymphoproliferative disease; autoimmune disease; hyperplasia; neurodegenerative disease; neoplasia; HIV; AIDS, cytostatic; human immunodeficiency virus; acquired immunodeficiency sylvis; telomere metabolism; anti-inflammatory; immunosuppressive; ss. New peptide nucleic acids that hybridizes to the RNA component of mammalian telomerase, useful for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or 0 Oligonucleotide #3 used in melting temperature studies of PNAs.

The present invention relates to peptide nucleic acids (PNAs), comprising

Example 2; Col 34; 46pp; English.

neurodegenerative diseases.

mammalian cells by hybridising to the RNA component of mammalian cells by hybridising to the RNA component of mammalian telomerase. The PNAs are useful as probes to detect the RNA component of mammalian telomerase and as inhibitors of telomerase activity, or to detect and/or quantitate polynucleotide having the human telomerase RNA component (hrR) sequence, as well as in forensic identification of individuals, such as paternity testing or identification of criminal suspects or unknown descendants based on the HTR gene RFLP pattern. The PNA can be further used for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases. The PNAs in combination with other pharmaceuticals (such as antineoplastic or cytostatic agents) can be used for treating neoplasia, hymen immunodeficiency virus (HIV) infections, acquired immunodeficiency syndrome (AIDS) and associated pathologies, and other diseases characterised by abnormal telomere metabolism or telomerase activity. The present sequence representing a DNA oligonucleotide is complementary to some of the PNAs of the present invention, and is used ö The invention relates to promoter regions from mouse and human telomerase RNA (TR) component genes. The TR gene promoter can be linked to a heterologous gene, especially a gene encoding a cytotoxin, for therapy of cancer, especially neoplasias. The telomerase is necessary for the unrestricted proliferative capacity of many human cancers. Mutation or dysregulation of the telomerase repression pathway may cause reactivation or upregulation of telomerase expression in cancer. Substances, and human telomerase RNA gene promoters, useful for tumor specific Gaps sequence of 6-25 nucleobases, that inhibit telomerase activity in Telomerase RNA; TR; promoter; cytotoxin; cancer; neoplasia; hTR; ; 0 3.5%; Score 16; DB 1; Length 16; 100.0%; Pred. No. 2.1e+02; tive 0; Mismatches 0; Indels Human telomerase RNA gene (hTR) specific primer hTRg. Sequence 16 BP; 5 A; 2 C; 6 G; 3 T; 0 U; 0 Other; (CANC-) CANCER RES CAMPAIGN TECHNOLOGY. Disclosure, Fig 6; 109pp; English. in melting temperature studies BP. 1 TAACTGAGAAGGGCGT 16 53 TAACTGAGAAGGGCGT 68 98GB-00001902. 99WO-GB000308. AAZ07277 standard; DNA; 20 (first entry) Query Match Best Local Similarity 100. Matches 16; Conservative WPI; 1999-479183/40. Homo sapiens 22-OCT-1999 WO9938964-A2 29-JAN-1999; 29-JAN-1998; PCR primer; 05-AUG-1999. gene therapy AAZ07277; Synthetic Keith WN; RESULT 286 Mouse \$5555555555555555555888 ò

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identified in the methods, can be used to block transcription from the TR gene promoter through interaction of the 5' regulatory sequences. These bubstances, e.g. antisense oligonucleotides, transcription factors, peptide nucleic acids and factors that disrupt signal transduction, are useful for cancer therapy. In particular, gene therapy vectors (especially pGT62-codAupp) comprising the promoter and a viral thymidine kinase gene can be used to convert a product, e.g. gancyclovir, so that neoplasia can be controlled or treated. Direct down-regulation of telomerase RNA gene through manipulation of transcription factors may be effective anticancer therapy and the cloning of the hTR gene promoter allows the analysis of therapeutic molecules which modulate hTR promoter activity. Sequences AAZ07621-80 represents PCR primers for amplifying human TR gene (HTR) promoter sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 function of nucleic acid molecules encoding hypotherical tumour endothelial marker. The antisense compound can also be used as research tools and diagnostics. It can also be used as tools in differential and/or combinatorial analyses to elucidate expression patterns of a portion or the entire complement of genes expressed within cells and tissues. The compound can also be used for treating diseases or conditions associated with hypothetical tumour endothelial marker, preferably hyperproliferative disorder, e.g. colon cancer. The compound can also be used as prophylaxis, e.g. to prevent or delay infection,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New antieense compound targeted to a nucleic acid molecule encoding hypothetical tumor endothelial marker, useful for modulating expression of hypothetical tumor endothelial marker or for treating colon cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to a compound targeted to a nucleic acid molecule encoding hypothetical tumour endothelial marker. The compound, particularly the antisense oligonucleotide is useful in modulating the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human tumour endothelial marker antisense oligonucleotide ISIS 208420.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            human; 88; antisense; hypothetical tumour endothelial marker; tumour; hyperproliferative disorder; colon cancer; infection; inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                  ;
0
                                                                                                                                                                                                                                                                                                                                                     3.5%; Score 16; DB 1; Length 20; 100.0%; Pred. No. 2.8e+02; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 4 A; 7 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 15; SEQ ID NO 71; 102pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                             451
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADH56499 Btandard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17-JUN-2002; 2002US-00174020.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 100."
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                             436 CTCGGCTCACACATGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Dobie KW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2004-061309/06.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-MAR-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADH56499;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADH56499/
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inflammation or tumour formation. The present sequence represents a human hypothetical tumour endothelial marker target region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to a compound targeted to a nucleic acid molecule particularly the antisense oligonucleotide is useful in modulating the function of nucleic acid molecules encoding hypothetical tumour endothelial marker. The compound, endothelial marker. The antisense compound can also be used as research tools and diagnostics. It can also be used as tools in differential and/or combinatorial analyses to elucidate expression patterns of a portion or the entire complement of genes expression patterns of a portion or the entire complement of genes expressed within cells and tissues. The compound can also be used for treating diseases or conditions associated with hypothetical tumour endothelial marker, preferably hyperproliferative disorder, e.g. colon cancer. The compound and also be used as prophylaxis, e.g. to prevent or delay infection, inflammation or tumour formation. The present sequence represents a human hypothetical tumour endothelial marker target region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New antisense compound targeted to a nucleic acid molecule encoding hypothetical tumor endothelial marker, useful for modulating expression of hypothetical tumor endothelial marker or for treating colon cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                    Human hypothetical tumour endothelial marker target region ISIS 126034.
                                                                                                                                       Gaрв
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           human; ss; hypothetical tumour endothelial marker; tumour;
hyperproliferative disorder; colon cancer; infection; inflammation.
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                                                                                              3.5%; Score 15.8; DB 1; Length 20; 89.5%; Pred. No. 2.9e+02;
                                                                                                                                       Indels
                                                       Sequence 20 BP; 1 A; 7 C; 4 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 8 A; 4 C; 7 G; 1 T; 0 U; 0 Other;
                                                                                                                                       0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 15; SEQ ID NO 138; 102pp; English.
                                                                                                                                                                          362 GGCCGCAGGAAGAACG 380
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    362 GGCCGCAGGAAGAGAACG 380
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           13
                                                                                                                                                                                                    GGCCACAGGAAAAGGAACG 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17-JUN-2002; 2002US-00174020.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17-JUN-2002; 2002US-00174020
                                                                                                                                                                                                                                                                                                                 ADH56566 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                Local Similarity 89.5
108 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Monia BP, Dobie KW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2004-061309/06.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18-DEC-2003.
                                                                                                                                                                                                                                                                                                                                                      ADH56566;
                                                                                                                                                                                                                 20
                                                                                                Query Match
                                                                                                                                       Matches
                                                                                                                                                                                                                                                                          RESULT 288
                                                                                                                                                                                                                                                                                                 ADH56566
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AAT89233;

RESULT 289

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This sequence is a novel peptide nucleic acid (PNA), which acts as an inhibitor of mammalian, preferably human, telomerase. The PNAs hybridise specifically to an RNA component of mammalian telomerase, and include the sequence GGG for specific hybridisation to the template region of this component. PNAs can be used as probes to detect the RNA component of mammalian telomerase and as inhibitors of telomerase activity, especially
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RBP-7; retinoblastoma binding protein-7; abnormal cell proliferation; diagnosis; therapy; cell differentiation; thyroid hyperplasia; psoriasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New peptide nucleic acids hybridising to mammalian telomerase RNA - used to inhibit telomerase, for treating tumours and other proliferative
                                                                                                                                                                                                                                                                             1..17
/*tag= a
//wod_base= OTHER
/node= "Sugar-phosphate backbone has been replaced by a
peptide backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Revised record issued on 21-OCT-2004 : Correction to feature table key
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PCR primer for human retinoblastoma binding protein-7 coding sequence.
                                                                                                                                                                  Peptide nucleic acid; PNA; cancer; telomerase; probe; hybridisation;
                                                                                                                            Peptide nucleic acid 9, targeted to mammalian telomerase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 17 BP; 2 A; 5 C; 4 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Piatyszek MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 9; Page 59; 76pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 diseases, also for diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      97WO-US005931.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 96US-00630019
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3.4%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17 AACCCTAACTGAGAGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAZ87074 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      in the treatment of cancer
                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16-MAY-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity 94.1
nes 16; Conservative
                                                             (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Wright WE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1997-512647/47.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (GERO-) GERON CORP.
                                                                                                                                                                                           .nhibitor; ss.
                                                                                                                                                                                                                                                                                 Key
modified base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        09-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   09-APR-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                  W09738013-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16-OCT-1997.
                                                             21-OCT-2004
                                                                                 12-MAY-1998
                                                                                                                                                                                                                                      Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Shay JW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 291
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This invention relates to a novel method concerning prognostic markers associated with EGFR (epidermal growth factor receptor) positive cancer. Specifically, it refers to a gene expression profiling method that can provide a prediction as to whether a patient is likely to respond well to treatment with an EGPR inhibitor. The present invention describes the cancer come gene selected from the group of CD4443, CD4446, DR5, GRO1, KRP17, IAMC2 or their products thereof. It further provides a cDNA microarray containing named genes that represent provides a cDNA corressing head or neck cancer or colon cancer exhibits elevated or decremaning whether a patient diagnosed with an EGFR expressing head or neck cancer or colon cancer exhibits elevated or decreased expression levels of these genes compared to normal. As such, these methods are also useful for prognosing or predicting the likelihood or cancer-attributable death or progression, including recurrence and metastatic spread of a neoplastic disease, as well as drug resistance. This polynucleotide sequence is a human PCR amplicon DNA sequence used as prognostic cancer marker, given in an exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Prognosing a patient with BGFR-expressing colon cancer comprises subjecting a sample comprising BGFR-expressing cancer cells to quantitative analysis of the expression level of the RNA transcript of at least one gene e.g., CD44v3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Gaps
                                                                                                                                                                                                                                             TERC; human; PCR amplicon; ds; prognostic marker; EGFR; epidermal growth factor receptor; cancer; gene expression profiling; microarray; head and neck cancer; colon cancer; metastatic spread; neoplastic disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 15.8; DB 1; Length 79;
Pred. No. 2.5e+02;
0; Mismatches 27; Indels
                                                                                                                                                                                                                    Human TERC DNA used as a cancer prognostic marker SeqID 84.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 79 BP; 16 A; 26 C; 27 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Shak S, Baselga J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 55; SEQ ID NO 84; 113pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENOMIC HEALTH INC. VALL HEBRON UNIV HOSPITAL.
                                                                                     ADP27647 standard; DNA; 79 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14-NOV-2003; 2003WO-US036777.
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Best Local Similarity 54.2%;
Matches 32; Conservative
                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cronin MT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2004-420643/39.
                                                                                                                                                                                                                                                                                                                                                                                                                  WO2004046386-A1.
                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                           26-AUG-2004
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                                                                                                                                ADP27647;
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Norton JC;

Corey D,

o;

Gaps

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Indels

Score 15.4; DB 1; Pred. No. 2.5e+02; 0; Mismatches 1;

64

BP.

AAT89233/c ID AAT89233 standard; DNA; 17 BP.

RESULT 290

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Length 17;

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14-AUG-2003; 2003WO-US025389.
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                                                                                                                                  ROSB SA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ROBB SA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 293
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  THE SECOND SECON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        요
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0
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                                                                                                                                                                                                                                                                                                                                                                                                                      novel nucleic acid and polymorphic markers used for diagnosis of diseases, especially those involving abnormal cell proliferation and differentiation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Acyl-coenzyme A synthetase 1, ACS1, antisense oligonucleotide #727.
benign prostate hypertrophy, cancer; sarcoma; neoplasm; leukaemia; lymphoma; biallelic marker; PCR primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                acyl-coenzyme A synthetase 1; ACS1; diabetes; obesity;
metabolic syndrome X; cardiovascular disorder; cancer; infection;
inflammation; tumour; antisense; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          and lymphomas. RBP-7 antibodies are useful as diagnostic agents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3.4%; Score 15.4; DB 1; Length 20; 94.1%; Pred. No. 3.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 9 A; 3 C; 4 G; 4 T; 0 U; 0 Other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 16; Page 206; 223pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   166 CAAACAAAAATGTCAG 182
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP.
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                                                                                                                                                                                                                                        98US-0091315P.
                                                                                                                                                                                               99WO-IB001242
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2000-117170/10.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO2004016749-A2.
                                                                                                           WO200000607-A1
                                                                                                                                                                                                                                                                                                           (GEST ) GENSET
                                                                                                                                                                                                                                                                                                                                                      Bougueleret L;
                                                                  Homo sapiens.
                                                                                                                                                                                               30-JUN-1999;
                                                                                                                                                                                                                                        30-JUN-1998;
10-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18-NOV-2004
                                                                                                                                                      06-JAN-2000.
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                                                                                                                                                                                                                                                                                                 New antisense compounds targeted to nucleic acid molecules encoding acyl-coenzyme A synthetase 1 (ACS1), useful for treating diseases or conditions associated with aberrant expression of ACS1, e.g. diabetes,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New antisense compounds targeted to nucleic acid molecules encoding acyl-coenzyme A synthetase 1 (ACS1), useful for treating diseases or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  acyl-coenzyme A synthetase 1; ACS1; diabetes; obesity;
metabolic syndrome X; cardiovascular disorder; cancer; infection;
inflammation; tumour; antisense; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 2 A; 13 C; 2 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 3; SEQ ID NO 727; 940pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             349
14-AUG-2002; 2002US-0403591P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 94.1
Matches 16; Conservative
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                                                                          (PHAA ) PHARMACIA CORP
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                                                                                                                                                                                                                               WPI; 2004-203782/19.
                                                                                                                                                                                                                                                                                                                                                                                                                       obesity or cancer
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US09436060A-16.rng.sl

Length 20;

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ADK21153 standard; DNA; 20
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              Local Similarity 85.0 nes 17; Conservative
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les 17; Conservative
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Query Match
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                       Best Loca
Matches
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Matches
                                                                                                                                                                                                       RESULT 295
                                                                                                                                                                                                                           ADK21153,
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                                                                                                        The invention relates to an antisense compound targeted to a nucleic acid molecule encoding acyl-coenzyme A synthetase 1 (ACS1). The antisense compound specifically hybridises with and inhibits the expression of ACS1. The antisense oligonuclectides or compounds are useful for inhibiting the expression of acyl-coenzyme A synthetase 1 (ACS1), and for treating diseases or conditions associated with aberrant expression of ACS1, e.g. diabetes, obesity, metabolic syndrome X, cardiovascular disorder or cancer. The antisense compounds are also useful as research reagents and kits, or in diagnostic, therapeutic and prophylactic applications, e.g. to prevent or delay infection, inflammation or tumour formation. The present sequence represents an acyl-coenzyme A synthetase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel rat pheochromocytoma PC12 cell line useful as cellular model of Huntington's disease, comprising an expression cassette containing a DNA encoding a truncated mutant huntingtin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention provides a rat or mouse pheochromocytoma cell line stably transfected with an expression cassette encoding a truncated mutant buntingtin. This can be used in the study of the pathological mechanism in Huntington's disease and enables the development of therapies and diagnostic techniques for the disease
conditions associated with aberrant expression of ACS1, e.g. diabetes,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      3.4%; Score 15.4; DB 1; Length 20;
4.1%; Pred. No. 3.1e+02;
ve 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 1 A; 12 C; 4 G; 3 T; 0 U; 0 Other;
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                                                                  Claim 3; SEQ ID NO 905; 940pp; English.
                                                                                                                                                                                                                                                                                                                                                                     1, ACS1, antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 1; Page 21; 55pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     332 CGGGGGCGAGGCGAGG 348
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CGAGGGCGAGGCCGAGG 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                94.1%;
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Best Local Similarity
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                          obesity or cancer.
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                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Acyl-coenzyme A synthetase 1, ACS1, antisense oligonucleotide #1230.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            acyl-coenzyme A synthetase 1; ACS1; diabetes; obesity;
metabolic syndrome X; cardiovascular disorder; cancer; infection;
inflammation; tumour; antisense; ss.
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                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
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3.4%; Score 15.2; DB 1;
85.0%; Pred. No. 3.2e+02;
iive 0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Mismatches
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                                                                                                                                             313 CTGTCAGCCGCGGGTCTCTC 332
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AAT89229/c
ID AAT89229 standard; DNA; 15 BP.
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(GERO-) GERON
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modified_base
                          WO9738013-A1
                                                               09-APR-1997;
                                                                                09-APR-1996;
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12-MAY-1998
                                            16-OCT-1997
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         Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic
                                                                                                                      Shay JW,
                                                                                                                                                                                                                                                                                                                                                                                                                                  AAT89226;
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                                                                                                                                                                                                                                                                                                                                                                         This sequence is a novel peptide nucleic acid (PNA), which acts as an inhibitor of mammalian, preferably human, telomerase. The PNAs hybridise specifically to an RNA component of mammalian telomerase, and include the sequence GGG for specific hybridisation to the template region of this component. PNAs can be used as probes to detect the RNA component of mammalian telomerase and as inhibitors of telomerase activity, especially in the treatment of cancer
                                                                                                                                                                                                                                                                                                                    New peptide nucleic acids hybridising to mammalian telomerase RNA - used to inhibit telomerase, for treating tumours and other proliferative diseases, also for diagnosis.
                                                                                                                                                         /note= "Sugar-phosphate backbone has been replaced by a peptide backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Revised record issued on 21-OCT-2004 : Correction to feature table key
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Peptide nucleic acid; PNA; cancer; telomerase; probe; hybridisation; inhibitor; human telomerase RNA; hTR; PCR; oligonucleotide; ss.
                                                                        Peptide nucleic acid; PNA; cancer; telomerase; probe; hybridisation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA oligonucleotide 4, used in the measurement of Tm values.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 15; DB 1; Length 15;
Pred. No. 2.3e+02;
                                                     Peptide nucleic acid 5, targeted to mammalian telomerase.
                                                                                                                                                                                                                                                                                 Norton JC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 15 BP; 5 A; 1 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                 Corey D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                 Piatyszek MA,
                                                                                                                     Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3.3.,
100.0%; Pre
                                                                                                                                                 mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                         Claim 9; Page 59; 76pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAT89248 standard; DNA; 15 BP.
                                                                                                                                                                                                                          97WO-US005931
                                                                                                                                                                                                                                            96US-00630019
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3.3%;
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                                    (first entry)
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Les 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15 TTGTCTAACCCTAAC
                           (revised)
                                                                                                                                                                                                                                                                                 Wright WE,
                                                                                                                                                                                                                                                                                                   WPI; 1997-512647/47.
                                                                                                                                                                                                                                                               (GERO-) GERON CORP.
                                                                                                                      Key
modified_base
                                                                                 inhibitor; ss
                                                                                                                                                                                      WO9738013-A1
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                                                                                                                                                                                                                                            09-APR-1996;
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                         21-OCT-2004
12-MAY-1998
                                                                                                                                                                                                       16-OCT-1997
                                                                                                  Synthetic
        AAT89229;
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                                                                                                                                                                                                                                                                                 Shay JW,
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                                                                                                                                                                                                                                                         New peptide nucleic acids hybridising to mammalian telomerase RNA - used to inhibit telomerase, for treating tumours and other proliferative diseases, also for diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          complimentary peptide nucleic acids (PNAs), (e.g. AAT89234). PNAs hybridise specifically to an RNA component of mammalian telomerase, and include the sequence GGG for specific hybridisation to the template region of this component. PNAs can be used as probes to detect the RNA component of mammalian telomerase and as inhibitors of telomerase activity, especially in the treatment of cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /mod_base= OTHER
/note= "Sugar-phosphate backbone has been replaced by a
peptide backbone"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Peptide nucleic acid; PNA; cancer; telomerase; probe; hybridisation; inhibitor; ss.
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0
                                                                                                                                                           Norton JC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3.3%; Score 15; DB 1; Length 15; 100.0%; Pred. No. 2.3e+02; ve 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 15 BP; 6 A; 4 C; 3 G; 2 T; 0 U; 0 Other;
                                                                                                                                                           Corey D,
                                                                                                                                                           Piatyszek MA,
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97WO-US005931.
                                                   96US-00630019
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ACCCTAACTGAGAAG 15
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                                                                                                                                                              Wright WE,
                                                                                                                                                                                                                 WPI; 1997-512647/47
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Best Local Similarity
Matches 15; Conserv
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oligonucleotides to the RNA component of human telomerase (hTR).
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                                                                                                                          This sequence is a novel peptide nucleic acid (DNA), which acts as an inhibitor of mammalian, preferably human, telomerase. The PNAs hybridise specifically to an RNA component of mammalian telomerase, and include the sequence GGG for specific hybridisation to the template region of this component. PNAs can be used as probes to detect the RNA component of mammalian telomerase and as inhibitors of telomerase activity, especially
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       - used for detecting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 for treating cancers, contraception,
                                                          New peptide nucleic acids hybridising to mammalian telomerase RNA - used to inhibit telomerase, for treating tumours and other proliferative diseases, also for diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        RNA component; human telomerase; antisense oligonucleotide; infection; neuroblastoma; bladder cancer; colon cancer; prostate cancer; cancer; contraception; sterilisation; immunosuppression; therapeutic; hTR; immune system down-regulation; anti-inflammatory therapy; ss.
                                                                                                                                                                                                                Revised record issued on 21-OCT-2004 : Correction to feature table key
                                                                                                                                                                                                                                                                              Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                     RNA component of human telomerase (hTR) antisense oligo 21ab3
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                      Corey D, Norton JC;
                                                                                                                                                                                                                                                        Length 15;
                                                                                                                                                                                                                                                                             0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New polynucleotide(s) anti:sense to human telomerase or inhibiting human telomerase, e.g. for treating car immuno-suppression or treating infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Weinrich SL;
                                                                                                                                                                                                                                   Seguence 15 BP; 3 A; 2 C; 5 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                        Score 15; DB 1; Lo
Pred. No. 2.3e+02;
                                                                                                                                                                                                                                                                              Mismatches
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                        Piatyszek MA,
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Local Similarity 100.0%; Pr
nes 15; Conservative 0;
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96US-00770565
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                                                                                                                                                                                             the treatment of cancer
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                                                                                                                                                                                                                                                                                                   CTAACCCTAACTGAG
                        Shay JW, Wright WE,
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                                         WPI; 1997-512647/47.
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    (GERO-) GERON CORP.
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                                                                                                                                                                                                                                                         Query Match
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                                                                                                                                                                                                                                                                                                                                                       RESULT 299
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continents of the many component of the hTR, but that does not sequence within an accessible region of the hTR, but that does not sequence within an accessible region of the hTR, but that does not component of thuman telomerase in a samplar region of hTR. These oligonucleotides may specifically be used for detection of an RNA component of human telomerase in a sample. This is useful for diagnosing cancer (especially neuroblastoma, bladder, colon and prostate cancer), and providing prognosis for a cancer patient. The inhibitory oligonucleotides can inhibit the telomerase activity level in a cell by interfering with transcription of the RNA component, decreasing the half-component into the telomerase holoenzyme, or inhibiting the polymerase activity of telomerase. These antisense oligonucleotides can be used for inhibiting telomerase. These antisense oligonucleotides can be used for inhibiting telomerase activity in both cultured cells and in cells in vivo. They can be used in therapeutics for treating or preventing cancer, for contraception or sterilisation, for immunosuppression, and for selectively down-regulating specific branches of inflammae system, e.g. a specific subset of T-cells, in anti-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Peptide nucleic acid; PNA; telomerase; ribonucleoprotein enzyme; cancer; inhibitor; neoplasia; neurodegenerative disease; aging; hyperplasia; AIDS; HIV; fungal infection; forensic identification; detect; tumour; paternity testing; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note= "Peptide nucleic acid molecule, where N-(2-aminoethyl)glycine units are linked to nucleotide bases via glycine amino N through a methylenecarbonyl linker"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
3.3%; Score 15; DB 1; Length 15;
Best Local Similarity 100.0%; Pred. No. 2.3e+02;
Matches 15; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 15 BP; 5 A; 3 C; 4 G; 3 T; 0 U; 0 Other;
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/*tag=
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Corey DR;

Norton JC,

Shay JW,

Piatyszek MA,

Wright WE,

WPI; 2000-292432/25

(TEXA) UNIV TEXAS SYSTEM.

96US-00630019.

09-APR-1996;

셤 ò The present sequence trepresents a peptide increase, and inhibits telomerase to the mRNA component of mammalian telomerase, and inhibits telomerase activity. Telomerase is a ribonucleoprotein enzyme that telomerase to synthesizes one strand of the telomeric DNA, using as a template and 11 nucleotide sequence contained within the RNA component of the enzyme. The invention relates to PNA molecules having a sequence of no more than 25 cases, which include the sequence GTTAGG. The uncharged nature of the PNA backbone increases the melting temperature of associating strands, increases the rate of association with targeted nucleic acids, and increases the rate of association with targeted nucleic acids, and therefore the requence of degradation by proceases or nucleases. The charapettic PNAs may be used for treating disease conditions such as cancers, neoplasia, hyperplasia, neurodegenerative diseases, aging, human conformation with antineoplastic and other cytcoxic or gyndrome) and associated pathologies, fungal infections, and other cytcoxic or cytcostatic agents, antifungal agents, and other nucleotides. PNAs may be used for molecular diagnostics, labelled PNAs are used as hybridization or individuals, e.g. paternity testing, based on hTR gene restriction of individuals, e.g. paternity testing, based on hTR gene restriction of individuals, e.g. paternity testing, based on hTR gene restriction of individuals, e.g. paternity testing, based on hTR gene restriction of individuals, e.g. paternity testing, based on hTR gene restriction of individuals, e.g. paternity testing, based on hTR gene and as a parterned and cancer and allows cancerous conditions to be detected with increase and as an earlow and earlows and earlows the paterned and and an earlow and allows and earlows conditions to be detected with increase and as an earlow the paterned and earlows the paterned and earlows conditions to be detected with increase and as an earlow and earlows. ö Peptide nucleic acid; PNA; telomerase; ribonucleoprotein enzyme; cancer; inhibitor; neoplasia; neurodegenerative disease; aging; hyperplasia; AIDS; HIV; fungal infection; forensic identification; detect; tumour; paternity testing; ss. possibly at an earlier stage, before cells are detected as cancerous based on pathological characteristics. The diagnostic and prognostic methods of the present invention can be used to detect an immortal or neoplastic cell or tumour tissue or cancer of any origin, provided the cell expresses telomerase activity and its RNA component The present sequence represents a peptide nucleic acid molecule which Gaps /*tag= a /note= "Phosphorothioate internucleotide linkages" ö Antisense sequence #45 used to inhibit telomerase activity. 3.3%; Score 15; DB 1; Length 15; 100.0%; Pred. No. 2.3e+02; ive 0; Mismatches 0; Indels Sequence 15 BP; 6 A; 4 C; 3 G; 2 T; 0 U; 0 Other; Location/Qualifiers Example 2; Col 33; 45pp; English AAA37587 standard; DNA; 15 BP 97US-00838545 63 1 ACCCTAACTGAGAAG 15 (first entry) 49 ACCCTAACTGAGAAG Conservative 1. .15 /*tag= Best Local Similarity Matches 15; Conserv 09-APR-1997; misc_feature 15-AUG-2000 US6046307-A. 04-APR-2000. Synthetic AAA37587; Query Match RESULT 301 셤 XXXCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCC

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The present sequence represents an antisense oligonucleotide used as a control sequence alongside a peptide nucleic acid molecule which control sequence alongside a peptide nucleic acid molecule which control sequence alongside a peptide nucleic acid molecule which telomerases activity. Telomerase is a ribonucleoprotein enzyme that telomerase activity. Telomerase is a ribonucleoprotein enzyme that controlled the sequence contained within the RNA component of the enzyme. The invention relates to PNA molecules having a sequence of no more than 25 bases, which include the sequence GTTAGG. The uncharged nature of the PNA concreases the melting temperature of associating strands, increases the melting temperature of associating strands.

CC diffords greater resistance of degradation by proteases or nucleases. The concerts, nooplassia, hyperplassa, neurodegenerative diseases. The mass cancers, nooplassia, hyperplassa, neurodegenerative diseases, aging, human immunodeficiency virus (HIV) infection/AIDS (acquired immunodeficiency syndrome) and associated pathologies, fungal infections, and other cyctoxic or syndrome) and associated pathologies, fungal infections, and other cyctoxic or cytostatic agents, antifungal agents, and other nucleotise. PNAs may be used for molecular diagnostics, labelled PNAs are used as hybridization probes to detect or quantitate polymorplasmic for forensic identification of individuals, e.g. paternity testing, based on hTR gene restriction fragment length polymorphism (RFLP) pattern. PNAs are also useful as cindwiduals, e.g. paternity testing, based on hTR gene restriction allows cancerous conditions to be detected with increased confidence and possibly at an earlier stage, before cells are detected as cancerous conditions to be detected with increased confidence and possibly at an earlier stage, before cells are detected as cancerous conditions to be detected with increased confidence and methods of the present invention can be used to detect an immorpary consisting and proposition allowanc
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New peptide nucleic acid (PNA) compounds that inhibit telomerase activilin mammalian cells is useful as probes to detect the RNA component of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3.3%; Score 15; DB 1; Length 15; 100.0%; Pred. No. 2.3e+02; ve 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 neoplastic cellor tumour tissue or cancer of any origin, cell expresses telomerase activity and its RNA component
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PNA sequence #2 used to inhibit telomerase activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 15 BP; 3 A; 2 C; 5 G; 5 T; 0 U; 0 Other;
                                                                                                                                                          Example 1; Col 27-28; 45pp; English.
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Best Local Similarity 100.0
Matches 15; Conservative
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                                                                                     mammalian telomerase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic
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AAA37545/c
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The present sequence to represents a pepticue increase, and inhibits to synthesizes to the mRNA component of mammalian telomerase, and inhibits telomerase activity. Telomerase is a ribonucleoprotein enzyme that telomerase synthesizes one strand of the telomeric DNA, component of the enzyme. The nucleotide sequence contained within the RNA component of the enzyme. The nucleotide sequence contained within the RNA component of the enzyme. The invention relates to PNA molecules having a sequence of no more than 25 bases, which include the sequence GTTAGG. The uncharged nature of the PNA backbone increases the melting temperature of associating strands, concers, the rate of association with targeted nucleic acids, and the rate of association with targeted nucleic acids, and the rate of association with targeted nucleic acids, and therefore the reapeutic PNAs may be used for treating disease or nucleases. The therapeutic PNAs may be used for treating disease conditions such as cancers, neoplasia, hyperplasia, neurodegenerative diseases, aging, human communication viths antineoplastic and other cytocoxic or diseases characterized by abnormal telomere metabolism or telomerase cytostatic agents, antifungal agents, and other nucleotides. PNAs may be used for molecular diagnostics, labelled PNAs are used as hybridization cytostatic agents, antifungal agents, and other nucleotides pained a human telomerase RNA (HTR) sequence. PNA probes are also used for forensic identification of individuals, e.g. paternity testing, based on hTR gene restriction fragment length polymorphism (RPLP) pattern. PNAs are also useful as probes to detect the RNA component of a mammalian telomerase and as inhibitors of telomerase activity. The method of the present invention allows cancerous conditions to be detected with increased conditions and content and content of the present invention allows cancerous conditions to be detected with increase and as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        wew peptide nucleic acid (PNA) compounds that inhibit telomerase activity in mammalian cells is useful as probes to detect the RNA component of a mammalian telomerase.
                                                                                           /note= "Peptide nucleic acid molecule, where N-(2-aminoethyl)glycine units are linked to nucleotide bases via glycine amino N through a methylenecarbonyl linker"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              possibly at an earlier stage, before cells are detected as cancerous based on pathological characteristics. The diagnostic and prognostic methods of the present invention can be used to detect an immortal or neoplastic cell or tumour tissue or cancer of any origin, provided the cell expresses telomerase activity and its RNA component
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            present sequence represents a peptide nucleic acid molecule which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Corey DR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Wright WE, Piatyszek MA, Shay JW, Norton JC,
                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 6; Col 71; 45pp; English.
                                                                                                                                                                                                                                                                                                                   97US-00838545.
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                        Key
misc_feature
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                         Gaps
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0
      Score 15; DB 1; Length 15;
Pred. No. 2.3e+02;
                        0; Indels
3.3%; Scor.
100.0%; Pred. No. 2...
0, Mismatches
                                             9
           Dest Local Similarity
watches 15; Conservative
                                             CTAACCCTAACTGAG
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        Query Match
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AAA37548 standard; DNA; 15 BP. 15 RESULT 303 AAA37548/ ID AAA3 셤

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15-AUG-2000 (first entry)
AAA37548;
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PNA sequence #5 used to inhibit telomerase activity.

Peptide nucleic acid; PNA; telomerase; ribonucleoprotein enzyme; cancer; inhibitor; neoplasia; neurodegenerative disease; aging; hyperplasia; AIDS; HIV; fungal infection; forensic identification; detect; tumour; paternity testing; ss.

Synthetic

/*tag= a //*tag= a "Peptide nucleic acid molecule, where N-(2-aminoethyl)glycine units are linked to nucleotide bases via glycine amino N through a methylenecarbonyl linker" Location/Qualifiers misc_feature

US6046307-A.

97US-00838545. 09-APR-1997;

96US-00630019 09-APR-1996;

(TEXA) UNIV TEXAS SYSTEM.

Corey DR; Wright WE, Piatyszek MA, Shay JW, Norton JC,

WPI; 2000-292432/25.

New peptide nucleic acid (PNA) compounds that inhibit telomerase activity in mammalian cells is useful as probes to detect the RNA component of a mammalian telomerase.

Claim 6; Col 71; 45pp; English.

The present sequence represents a peptide nucleic acid molecule which hybridises to the mRNA component of mammalian telomerase, and inhibits call chowerase activity. Telomerase is a ribonucleoprotein enzyme that synthesizes one strand of the telomeric DNA, using as a template an 11 nucleotide sequence contained within the RNA component of the enzyme. The nucleotide sequence contained within the RNA component of the enzyme. The nucleotide sequence contained within the RNA component of the enzyme. The nucleotide sequence contained within the RNA component of the enzyme. The asses, which include the sequence GTTAGG The uncharged nature of the PNA backbone increases the melting temperature of associating strands, cancers season the rate of association with targeted nucleic acids, and affords greater resistance of degradation by proteases or nucleases. The therapeutic PNAs may be used for treating disease conditions such as cancers neoplasia, hyperplasia, neurodegenerative diseases. The characterized by abnormal telomerative diseases, and other cytostatic agents, antifungal infection, and other volencies. Full of the activity, in combination with antineoplastic and other cytostacic agents, antifungal agents, and other nucleotides. PNAs may be used for molecular diagnostics, labelled PNAs are used as hybridization of cytostatic agents, antifungal agents, and other nucleotides. PNAs may be used for molecular diagnostics, labelled PNAs are used as hybridization of individuals, e.g. paternity testing, based on hTR gene restriction of individuals, e.g. paternity testing, based on hTR gene restriction of individuals, e.g. paternity testing, based on hTR gene serviction of an anilows cancerous conditions that the RNA component of a mammalian telomerase and as allows cancerous conditions that the patern of a defect of a defect the RNA component of a mammalian defence and a sallows cancerous conditions to be detected with increased conditions. based on pathological characteristics. The diagnostic and prognostic methods of the present invention can be used to detect an immortal or neoplastic cell or tumour tissue or cancer of any origin, provided the possibly at an earlier stage, before cells are detected as cancerous cell expresses telomerase activity and its RNA component

Sequence 15 BP; 5 A; 1 C; 5 G; 4 T; 0 U; 0 Other;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention relates to peptide nucleic acids (PNAs), comprising a sequence of 6-25 nucleobases, that inhibit telomerase activity in mammaalian cells by hybridising to the RNA component of mammalian telomerase and as inhibitors of telomerase activity, or to detect and/or quantitate polynucleotide having the human telomerase RNA component (HTR) sequence, as well as in forensic identification of individuals, such as paternity testing or identification of criminal suspects or unknown descendants based on the HTR gene RFLP pattern. The PNA can be further used for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases. The PNAs in combination with other pharmaceuticals (such as antineoplastic or cytostatic agents) can be used for treating neoplasia,
                                                                                                                                                                                                                                                                                                                                                              /*tag= a
/note= "This sequence is a peptide nucleic acid, i.e. it
contains a polyamide backbone instead of a deoxyribose
backbone"
                                 Gaps
                                                                                                                                                                                                                       Mammalian; peptide nucleic acid; probe; forensic; paternity testing; human telomerase RNA component; HTR gene RFLP pattern; cancer; inflammation; lymphoproliferative disease; autoimmune disease; neurodegenerative disease; neoplasia; hyperplasia; HIV; AIDS; human immunodeficiency virus; acquired immunodeficiency syndrome; telomere metabolism; mutant; cytostatic; anti-inflammatory; immunosuppressive; polyamide backbone; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New peptide nucleic acids that hybridizes to the RNA component of mammalian telomerase, useful for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases.
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                                                                                                                                                                                                     PNA XIII inhibiting human and mammalian telomerase activity.
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            Length 15;
                                 0; Indels
           Score 15; DB 1; Le
Pred. No. 2.3e+02;
                               Mismatches
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    3.3%; Scc.
100.0%; Pred
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97US-00838545.
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                                                                                                                          427/c
AAS15427 standard; DNA; 15
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                                                                                                                                                                              (first entry)
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                                                       TTGTCTAACCCTAAC
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Query Match
Best Local Similarity
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09-APR-1997;
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                                                                                                                                                                                                                                                                                                                              Synthetic.
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The present invention relates to peptide nucleic acids (PNAs), comprising a sequence of 6-25 nucleobases, that inhibit telomerase activity in mammalian cells by hybridising to the RNA component of mammalian telomerase and as inhibitors of telomerase activity, or to detect and/or quantitate polynucleotide having the human telomerase RNA component (hTR) sequence, as well as in forensic identification of individuals, such as paternity testing or identification of criminal suspects or unknown descendants based on the hTR gene RFLP pattern. The PNA can be further used for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases. The PNAs in combination with other pharmaceuticals (such as antinooplastic or cycostatic agents) can be used for treating neoplasia, hymen immunodeficiency virus (HIV) infections, acquired
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                                                                                                   activity. The present sequence represents one of the PNA sequences of the
hyperplasia, human immunodeficiency virus (HIV) infections, acquired immunodeficiency syndrome (AIDS) and associated pathologies, and other diseases characterised by abnormal telomere metabolism or telomerase
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Oligonucleotide #4 used in melting temperature studies of PNAs.
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                                                                                                                                                                                                                                                           Length 15;
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                                                                                                                                                                                            G; 4 T; 0 U; 0 Other;
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2.3e+02;
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                                                                                                                                                                                                                                                    3.3%; Score 15; DB 100.0%; Pred. No. 2.3 ative 0; Mismatches
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97US-00838545
                                                                                                                                                                                               Sequence 15 BP; 5 A; 1 C; 5
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Best Local Similarity
Matches 15; Conserv
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09-APR-1997;
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                                                                                                                                       invention
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RESULT 307
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/note= "This sequence is a peptide nucleic acid, i.e. it
contains a polyamide backbone instead of a deoxyribose
backbone"
             diseases characterised by abnormal telomere metabolism or telomerase activity. The present sequence representing a DNA oligonucleotide is complementary to some of the PNAs of the present invention, and is used
 immunodeficiency syndrome (AIDS) and associated pathologies, and other
                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                        Mammalian; peptide nucleic acid; probe; forensic; paternity testing; human telomerase RNA component; hTR gene RFLP pattern; cancer; inflammation; lymphoproliferative disease; autoimmune disease; neurodegenerative disease; neoplasia; hyperplasia; HIV; AIDS; human immunodeficiency virus; acquired immunodeficiency syndrome; telomere metabolism; mutant; cytostatic; anti-inflammatory; immunosuppressive; polyamide backbone; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New peptide nucleic acids that hybridizes to the RNA component of mammalian telomerase, useful for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, on neurodegenerative diseases.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Norton JC;
                                                                                          Length 15;
                                                                                                               Indels
                                                                 Sequence 15 BP; 6 A; 4 C; 3 G; 2 T; 0 U; 0 Other;
                                                                                        3.3%; Score 15; DB 1; Le
100.0%; Pred. No. 2.3e+02;
ive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
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                                              melting temperature studies
                                                                                                                                                                                                                   ВР
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                                                                                                                                     ACCCTAACTGAGAAG 63
                                                                                                                                                           1 Acceraácicadad 15
                                                                                                                                                                                                                   AAS15424 standard; DNA; 15
                                                                                                                                                                                                                                                               (first entry)
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                                                                                                    Local Similarity 100.
nes 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Shay JW, Wright WE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2001-638024/73.
                                                                                                                                                                                                                                                                                                                                                                                                                                               modified base
                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US6294650-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              08-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    09-APR-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              09-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
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                                                                                           Query Match
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                                                                                                                                                                                                                                                                                                      activity. The present sequence represents one of the PNA sequences of the
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     component;
                                                                                                                     lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases. The PNAs in combination with other pharmaceuticals (such as antineoplastic or cytostatic agents) can be used for treating neoplasia, hyperplasia, human immunodeficiency virus (HIV) infections, acquired immunodeficiency syndrome (AIDS) and associated pathologies, and other diseases characterised by abnormal telomere metabolism or telomerase.
                                                                                     PNA can be further used for treating or preventing cancer, inflammation,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mammalian; forensic; paternity testing; human telomerase RNA component; hTR gene RFLP pattern; cancer; inflammation; lymphoproliferative diseas autoimmune disease; neurodegenerative disease; neoplasia; hyperplasia; HIV; AIDS; human immunodeficiency virus; telomere metabolism; mutant; acquired immunodeficiency syndrome; cytostatic; anti-inflammatory; immunosuppressive; phosphorothloate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Phosphorothioate (PS) oligomer II used to inhibit telomerase activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
component (hTR) sequence, as well as in forensic identification of individuals, such as paternity testing or identification of criminal suspects or unknown descendants based on the hTR gene RFLP pattern.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New peptide nucleic acids that hybridizes to the RNA component of mammalian telomerase, useful for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note= "Phosphorothioate internucleotide linkages"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 15; DB 1; Length 15;
Pred. No. 2.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 15 BP; 3 A; 2 C; 5 G; 5 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                        3.3%; Scc...
100.0%; Pred. No. ...
0; Mismatches
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OTHER
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97US-00838545.
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Best Local Similarity 100.
Matches 15; Conservative
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modified_base
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                                                                                                                                                                                                                                                                                                                                                  nvention
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a sequence of 6-25 nucleobases, that inhibit telomerase activity in mammalian cells by hybridising to the RNA component of mammalian telomerase. The PMAs are useful as probes to detect the RNA component of mammalian telomerase and as inhibitors of telomerase activity, or to
                                                                  detect and/or quantitate polymicleotide having the human telomerase RNA component (HTR) sequence, as well as in forensic identification of individuals, such as paternity testing or identification of criminal suspects or unknown descendants based on the hTR gene RFLP pattern. The PNA can be further used for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases. The PNAs in combination with other pharmaceuticals (such as antinoeplastic or cytostatic agents) can be used for treating neoplasta human immunodeficiency virus (HIV) infections, acquired immunodeficiency syndrome (AIDS) and associated pathologies, and other diseases characterised by abnormal telomere metabolism or telomerase activity. The present sequence represents a phosphorothicate (PS) oligomer used to inhibit telomerase activity in the methods of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New polynucleotide useful for inhibiting telomerase activity in cells, c for treating telomerase-mediated condition or disease, such as cancers, tumors, Hodgkin's disease, or inflammatory conditions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; telomerase; hTR; cytostatic; anti-inflammatory; adenocarcinoma;
breast; prostate; colon; mixed cell leukaemia; Hodgkin's disease;
fertility; inflammatory condition; tumour; cancer; veterinary;
immunosuppression; telomerase inhibitor; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                 3.3%; Score 15; DB 1; Length 15;
100.0%; Pred. No. 2.3e+02;
rative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                  Seguence 15 BP; 3 A; 2 C; 5 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human telomerase polynucleotide inhibitor #8.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Weinrich SL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
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/*tag= a
/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 8; Page 36; 48pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                             46 CTAACCCTAACTGAG 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15 criadcccriadcreag 1
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Best Local Similarity 100.
                                                                                                                                                                                                                                                                                                                present invention
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The invention relates to polynucleotide inhibitors (I) and methods for inhibiting telomerase activity. (I) are useful in inhibiting telomerase activity and proliferation of a telomerase positive cell, and in manufacturing a medicament for inhibiting telomerase activity in a class and in treating telomerase-mediated condition or disease, such as adenocarcinoma of breast, proctate or colon, mixed cell leukaemia, Hodgkin's disease, fertility and inflammatory conditions. (I) are also useful in treating a tumour or in manufacturing a medicament for the treatment of tumour. The polynucleotide inhibitors may also be used in diagnostic assays for detecting RNA or DNA. Inhibitors and therapeutic methods of treating cancer and other disorders involving inappropriate expression of telomerase, and in treating veterinary proliferative captesion of telomerase, and in treating veterinary proliferative captession of telomerase in hammatopoietic stem cells is useful for immunosuppression and for selectively down-regulating specific for immunosuppression and for selectively down-regulating specific contended of the immune system. The present sequence represents human telomerase polynucleotide inhibitor #8, as described in the method of the
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     for treating telomerase-mediated condition or disease, such as cancers, tumors, Hodgkin's disease, or inflammatory conditions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; telomerase; hTR; cytostatic; anti-inflammatory; adenocarcinoma; breast; prostate; colon; mixed cell leukaemia; Hodgkin's disease; fertility; inflammatory condition; tumour; cancer; veterinary; immunosuppression; telomerase inhibitor; ss.
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100.0%; Pred. No. 2.3e+02;
rative 0; Mismatches 0; Indels
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/note= "N3'-P5' phosphoramidate linkages"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 15 BP; 3 A; 2 C; 9 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human telomerase polynucleotide inhibitor #12
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AAS15931 standard; DNA; 15
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Best Local Similarity
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XX AAS3

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                        The invention relates to polymucleotide inhibitors (I) and methods for inhibiting telomerase activity. (I) are useful in inhibiting telomerase activity and proliferation of a telomerase positive cell, and in manufacturing a medicament for inhibiting telomerase activity in a cell and in treating telomerase-mediated condition or disease, such as adenocarcinoma of breast, prostate or colon, mixed cell leukaemia, CK Hodgkin's disease, fertility and inflammatory conditions. (I) are also useful in treating a tumour or in manufacturing a medicament for the ctreatment of tumour. The polymucleotide inhibitors may also be used in catament of tumour. The polymucleotide inhibitors may also be used in catament of tumour or in useful in prophylactic and therapeutic activity in cells in vivo is useful in prophylactic and therapeutic methods of treating cancer and other disorders involving inappropriate expression of telomerase, and in treating veterinary proliferative configences in thibition of telomerase in haematopoletic stem cells is useful for immunosuppression and for selectively down-regulating specific branches of the immune system. The present sequence represents human the immonerace polynucleotide inhibitor #12, as described in the method of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                'note= "N3'-P5' phosphoramidate linkages"
                                                                                                                                                                                                                                                                                                                     Sequence 15 BP; 6 A; 1 C; 6 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human telomerase polynucleotide inhibitor #13.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
Example 3; Page 32; 48pp; English.
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/mod_base= OTHER
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AAS15932 standard; DNA; 15 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (GERO-) GERON CORP.
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                                                                                                                                                                                                                                                                                             the invention
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modified_base
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New polynucleotide useful for inhibiting telomerase activity in cells, or for treating telomerase-mediated condition or disease, such as cancers, tumors, Hodgkin's disease, or inflammatory conditions.

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The invention relates to polymucleotide inhibitors (I) and methods for inhibiting telomerase activity. (I) are useful in inhibiting telomerase activity. (I) are useful in inhibiting telomerase activity and proliferation of a telomerase positive cell, and in manufacturing a medicament for inhibiting telomerase activity in a cell and in treating telomerase-mediated condition or disease, such as adenocarcinoma of breast, prostate or colon, mixed cell leukaemia. (Consection of disease, fertility and inflammatory conditions. (I) are also treatment of tumour or in manufacturing a medicament for the treatment of tumour. The polymucleotide inhibitors may also be used in diagnostic assays for detecting RNA or DNA. Inhibition of telomerase cation cells in vivo is useful in prophylactic and therapeutic methods of treating cancer and other disorders involving inappropriate expression of telomerase, and in treating veterinary proliferative capression of telomerase, and in treating veterinary proliferative for immunosuppression and for selectively down-regulating specific branches of the immune system. The present sequence represents human the inversion that in the present sequence represents human the inversion.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3.3%; Score 15; DB 1; Length 15; 100.0%; Pred. No. 2.3e+02; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 15 BP; 5 A; 1 C; 7 G; 2 T; 0 U; 0 Other;
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/mod_base= OTHER
/note= "A-hydroxyethyl phosphate"
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Example 3; Page 32; 48pp; English.
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ID ADP87875 standard; DNA; 15
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Best Local Similarity 100.
Matches 15, Conservative
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12-OCT-1998
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   therapy.
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                                                                                                              The present invention relates to novel 2',5'-oligoadenylic acid analogs and their pharmacologically- acceptable sakies. The analogs are stable with superior antitumour and antiviral activity and so are useful in cancer or antiviral therapy e.g. as antisense drugs. The present sequence
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nucleic acid units as well as unusual phosphate groups with excellent activity particularly antitumor, applicable in cancer or antiviral
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11. .15
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/note= "A-hydroxyethyl phosphate"
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                                                                            Disclosure; Page 100; 220pp; Japanese
                                                                                                                                                                                         was used to illustrate the invention.
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/mod_base= OTHER
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Best Local Similarity
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                                         therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 312
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                                                                   The present invention relates to novel 2',5'-oligoadenylic acid analogs and their pharmacologically- acceptable salts. The analogs are stable with superior antitumour and antiviral activity and so are useful in cancer or antiviral therapy e.g. as antisense drugs. The present sequence was used to illustrate the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PP2; human; telomerase protein 2; cancer; AIDS; ageing; therapy; PCR;
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                                                                                                                                                                                                                                                                                                                                                                           3.3%; Score 15; DB 1; Length 15; 100.0%; Pred. No. 2.3e+02; ve 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                   Sequence 15 BP; 5 A; 3 C; 7 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human telomerase antisense RNA primer.
Disclosure; Page 100; 220pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 8; Page 91; 150pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ВР
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         96US-00751189.
97US-00873039.
97US-00951733.
                                                                                                                                                                                                                                                                                                                                                                                                                       100.08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     97WO-US021248
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Harrington LA, Robinson MO;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       80 TTTTGCTCCCGCGC 94
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                immune deficiency syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TTTTGCTCCCGCGC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAV27891 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INC.
CANADA INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1998-297946/26
                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
nes 15; Conserv
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ADK20144/
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to an antisense compound targeted to a nucleic acid molecule encoding acyl-coenzyme A synthetase 1 (ACS1). The antisense compound specifically hybridises with and inhibits the expression of ACS1. The antisense oligonucleotides or compounds are useful for inhibiting the expression of acyl-coenzyme A synthetase 1 (ACS1), and for treating diseases or conditions associated with aberrant expression of ACS1, e.g. diabetes, obesity, metabolic syndrome X, cardiovascular reagents and kite, or in diagnostic, therapeutic and prophylactic applications, e.g. to prevent or delay infection, inflammation or tumour formation. The present sequence represents an acyl-coenzyme A synthetase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New antisense compounds targeted to nucleic acid molecules encoding acyl-coenzyme A synthetase 1 (ACS1), useful for treating diseases or conditions associated with aberrant expression of ACS1, e.g. diabetes,
                                                                          Gaps
                                                                                                                                                                                                                                              Acyl-coenzyme A synthetase 1, ACS1, antisense oligonucleotide #205.
                                                                                                                                                                                                                                                                     acyl-coenzyme A synthetase 1; ACS1; diabetes; obesity;
metabolic syndrome X; cardiovascular disorder; cancer; infection;
inflammation; tumour; antisense; ss.
                                                                         .;
0
                                             Score 15; DB 1; Length 18;
Pred. No. 2.9e+02;
0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3.3%; Score 15; DB 1; Length 20; 100.0%; Pred. No. 3.38+02; ive 0; Mismatches 0; Indels
                     Sequence 18 BP; 1 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 3 A; 12 C; 2 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 3; SEQ ID NO 205; 940pp; English.

    ACS1, antisense oligonucleotide.

                                            Query Match 3.3%; Sco
Best Local Similarity 100.0%; Pr
Matches 15; Conservative 0;
                                                                                                                                                                          ADK20128 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                        14-AUG-2003; 2003WO-US025389.
                                                                                                                                                                                                                                                                                                                                                                                                                14-AUG-2002; 2002US-0403591P.
                                                                                                         GGGTGGGCCTGGGAG 18
                                                                                           11 GGGTGGGCCTGGGAG 25
                                                                                                                                                                                                                        18-NOV-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity 100.
nes 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                    (PHAA ) PHARMACIA CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2004-203782/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             obesity or cancer.
                                                                                                                                                                                                                                                                                                                                          WO2004016749-A2.
                                                                                                                                                                                                                                                                                                                                                                 26-FEB-2004
                                                                                                                                                                                                                                                                                                                     Synthetic.
                                                                                                                                                                                                  ADK20128;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ross SA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
field.)
                                                                                                                                                    RESULT 314
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Loca
Matches
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The invention relates to an antisense compound targeted to a nucleic acid molecule encoding acyl-coenzyme A synthetase 1 (ACS1). The antisense compound specifically hybridises with and inhibits the expression of ACS1. The antisense oligonucleotides or compounds are useful for inhibiting the expression of acyl-coenzyme A synthetase 1 (ACS1), and for treating diseases or conditions associated with aberrant expression of ACS1, e.g. diabetes, obesity, metabolic syndrome X, cardiovascular disorder or cancer. The antisense compounds are also useful as research reagents and kits, or in diagnostic, therapeutic and prophylactic applications, e.g. to prevent or delay infection, inflammation or tumour formation. The present sequence represents an acyl-coenzyme A synthetase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New antisense compounds targeted to nucleic acid molecules encoding acylcoenzyme A synthetase 1 (ACS1), useful for treating diseases or conditions associated with aberrant expression of ACS1, e.g. diabetes,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cold-adapted influenza virus; passage culture; PB2 protein; PB1 protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                   Acyl-coenzyme A synthetase 1, ACS1, antisense oligonucleotide #221.
                                                                                                                                                                                                                        acyl-coenzyme A synthetase 1; AC31; diabetes; obesity;
metabolic syndrome X; cardiovascular disorder; cancer; infection;
inflammation; tumour; antisense; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3.3%; Score 15; DB 1; Length 20; 100.0%; Pred. No. 3.3e+02; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 3 A; 13 C; 2 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Influenza virus PA gene specific primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 3; SEQ ID NO 221; 940pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1, ACS1, antisense oligonucleotide
ADK20144 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14-AUG-2003; 2003WO-US025389.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            14-AUG-2002; 2002US-0403591P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ВР
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                335 GGGCGAGGCGAGGT 349
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Beet Local Similarity 100.00
Conservative
Conservative
                                                                                                            18-NOV-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAX82232 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20 GGCGAGGCCGAGGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (PHAA ) PHARMACIA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2004-203782/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     obesity or cancer.
                                                                                                                                                                                                                                                                                                                                                                                           WO2004016749-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   26-FEB-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       18-AUG-1999
                                                                                                                                                                                                                                                                                                                                      Synthetic.
                                                       ADK20144;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ross SA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAX82232;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 316
AAX82232/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           M X G X G X G X G
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Gaps

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GGGCGAGGCGAGGT 349

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19 GGGCGAGGCGAGGT 5

99WO-US008731.

98US-0082477P

Reid TM;

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A novel rapid and sensitive method using PCR for detecting Histoplasma
                                                                              (USSH ) US DEPT HEALTH & HUMAN SERVICES.
                                                                                                                                                           capsulatum in samples.
                                                                                                                         MPI; 2000-052703/04.
                                 20-APR-1999;
                                                       21-APR-1998;
                                                                                                    Schafer MP,
           28-OCT-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAQ23265;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     à
                                                                                                                                                                                                                                                                                  The invention relates to cold-adapted influenza viruses prepared by passage culture of A/X-31, B/Yamagata/16/88 or B/Lee/40 viruses at low temperatures A cDNA gene of cold-adapted influenza virus HTCA-A101 can be selected from a group consisting of PB2 protein gene, PB1 protein gene, RAPRS129.2 x813197). The method is useful for the production of coldadapted influenza virus that exhibit temperature sensitivity and can be actively grown in fertilized eggs. The virus is useful for vaccines for protection against flu. Live vaccines containing cold-adapted viruses have several advantages over killed vaccines. It can prevent reduction of immunogenicity, which may occur in the killed vaccine where antigenic protecins would be denatured at its inactivation. It can also avoid proteins it promotes the immunity by inducing IgA and it can be administered into a spray formulation via nasal cavity and thus its administered into a spray formulation via nasal cavity and thus its convenient for children. It she expected.

Sequences AXX82222-X82257 represent PCR primers specific for the various
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
PA protein; NP protein; M protein; NS protein; temperature sensitivity; vaccine; flu; influenza; PCR primer; ss.
                                                                                                                                                                                                                            Cold-adapted influenza viruses useful for the production of protective
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                  Kim HG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        5.8S rRNA; detection; environmental sample; public health; soil; vegetation; dust; faeces; histoplasmosis; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 14.8; DB 1; Length 18; Pred. No. 3e+02; 0; Mismatches 2; Indels
                                                                                                                                                                                  Kim J,
                                                                                                                                                                                  Cheoun KH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 18 BP; 8 A; 4 C; 5 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    H. capsulatum 5.8S rRNA gene PCR primer HC2.
                                                                                                                                                                                  Kim SJ,
                                                                                                                                                                                                                                                                Example 4; Page 15; 62pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        29 TGGTGGCCATTTTTTGTC 46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18 recreecearricare 1
                                                                                                               98WO-KR000384.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3.3%;
88.9%;
                                                                                                                                                                                  Youn JW,
                                                                                                                                      97KR-00064854
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAZ24498 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (revised)
(first entry)
                                                                                                                                                            (CHEI-) CHEIL JEDANG CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 88.9*
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              genes of influenza virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ajellomyces capsulatus
                                                                                                                                                                                                                                           vaccines against flu.
                                                                                                                                                                                  Lee KH,
                                                                                                                                                                                                       WPI; 1999-385377/32.
                                              Influenza virus.
                                                                                                               30-NOV-1998;
                                                                                                                                      29-NOV-1997;
                                                                  WO9928445-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15-SEP-2003
18-FEB-2000
                                                                                         10-JUN-1999
                                  Synthetic.
                                                                                                                                                                                   Seong BL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAZ24498;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 317
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAZ24498/c
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This invention describes a novel rapid and sensitive method for detecting Histoplasma capsulatum. The method uses oligonucleotide primers which amplify a segment of DNA specific to the H. capsulatum 5.85 rRNA gene. The method of the invention can be used to detect H. capsulatum in a wide variety of samples. For example, for public health protection convironmental samples containing soil, vegetation, dust, decaying faeces for midial packs etc. can be assayed for the presence of the canalysed e.g. those who are suspected of having contracted fresh histoplasmosis as a result of symptomology, and immune compromised individuals who may be at risk of contracting orhonic pulmonary or ocular chaptasmosis. The prior arisk of contracting orhonic pulmonary or ocular chaptasmosis. The prior arisk of contracting orhonic pulmonary or capsulatum is expensive and requires several weeks to complete. The capsulatum is expensive and requires several weeks to complete. The sample are collected, small but highly contaminated areas can be overlooked. A need exists for a specific, sensitive, and rapid assay for H. capsulatum. The present invention provides this need, as it is rapid, sensitive and inexpensive. This sequence represents PCR primer HCI which is used in the method of detection. (Updated on 15-SEP-2003 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 18 BP; 3 A; 8 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Japanese type C hepatitis virus partial gene.
Claim 2; Page 30; 33pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      410 CTGAGCTGTGGGACGTGC 427
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          90JP-00154204.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAQ23265 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HCV; probe; diagnostic; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              23-JUL-1992 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (TOKU ) TOKUYAMA SODA KK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity 88.9
nes 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hepatitis C virus
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Freier SM;
inhibitor
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                                                                                                                                                                                    Query Match
                                                                                                                                                                                                 Best Loca
Matches
                                                                                                                                                                                                                                                                                                RESULT 320
                                                                                                                                                                                                                                                                                                             ADH62909
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                                                                                                        The polymucleotide contains the partial base sequence of the Japanese type C hepatitis virus (HCV) gene. The DNA was obtd. from the serum of a non A non B hepatitis patient. The serum was ultracentrifuged and the ppte. dissolved in a mixture of GITC, Na citrate, sarcosine and mercaptoethanol. Phenol chloroform was added and the mixture centrifuged. Isopropanol was added to the aq. layer which was left to stand at -20 deg C for 3 hrs, then centrifuged. The ppte. was treated by GITC and phenolchloroform and pptd. by ethanol and dissolved in di20 to give an aq. DNA soln. The gene was amplified by PCR and the HCV gene cloned for determination of its base sequence. A genomic version of the sequence was given (AAQ23036). In the version shown there is a Y at base 189. The DNA in the specification has a guanine at this position. The DNA may be used as a probe for detection of Japanese type C hepatitis. See also AAQ23264
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention describes a method for inducing apoptosis in a cell or animal comprising administering to a cell or animal a Jagged 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         apoptosis; Jagged 2 inhibitor; cytostatic; hyperproliferative disorder; human; ss; PCR primer.
                                     Poly:nucleotide for hepatitis virus gene detection - using poly:nucleotide deoxyribonucleic acid probe, hybridising with hepatitis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Inducing apoptosis in a cell or animal for treating a subject having a condition associated with insufficient apoptosis by administering to a cell or animal a Jagged 2 inhibitor to reduce Jagged 2 levels or
                                                                                                                                                                                                                                                                                                                                      Gaps
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                                                                                                                                                                                                                                                                                                            3.2%; Score 14.6; DB 1; Length 18; 82.4%; Pred. No. 3.18+02; tive 2; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                     Sequence 18 BP; 1 A; 10 C; 5 G; 0 T; 0 U; 2 Other;
                                                poly:nucleotide deoxyribonucleic acid probe, h
virus gene and detecting hybrid for diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human Jagged 2 forward PCR primer SEQ ID NO:4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 13; SEQ ID NO 4; 148pp; English
                                                                                     Disclosure; Page 2; 14pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                              196 CGCCCTCCCGGGGACC 212
                                                                                                                                                                                                                                                                                                                                                                           *GCCCCCCYGGGGACC 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10-MAR-2003; 2003WO-US007340
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12-MAR-2002; 2002US-00096399
                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADD00949 standard; DNA; 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                      14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Shapard PJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-756943/71.
              WPI; 1992-101941/13.
                                                                                                                                                                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO2003077848-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Koller E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADD00949;
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                                                                                                                                                                                                                                                                                                             Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Antisense, Jagged 2; hyperproliferative disorder; cancer; developmental disorder; apoptosis; prophylaxis; antisense therapy; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New antisense oligonucleotides for modulating Jagged 2 expression, useful for diagnosing, preventing or treating diseases or conditions associated with Jagged 2, e.g. cancer or developmental disorders.
                    treating a subject having a disease or condition associated with insufficient apoptosis by administration of a Jagged 2 inhibitor; (2) a pharmaceutical composition comprising a Jagged 2 inhibitor and another active ingredient for inducing apoptosis; and (3) a kit comprising a Jagged 2 inhibitor and instructions for using the Jagged 2 inhibitor in the induction of apoptosis. The Jagged 2 inhibitor has cytostatic activity. The method can be used for inducing apoptosis in a cell or animal for treating a subject having a disease or condition associated with insufficient apoptosis, e.g., hyperproliferative disorder. The present sequence represents a PCR primer for human Jagged 2, which is used in an example from the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
to reduce Jagged 2 levels or activity. Also described:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3.2%; Score 14.4; DB 1; 93.8%; Pred. No. 2.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human Jagged 2 DNA specific forward PCR primer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               265 CCCGGGGCTTCTCCGG 280
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hes 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-898250/82.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           primer; PCR; ss.
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NFKB sub-unit modulating inozyme substrate #146.

(first entry)

03-JUN-2003

ACA06327;

BP.

ACA06327 standard; RNA; 17

ACA06327 ID ACA(

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This invention relates to novel antisense compounds that can be used to modulate the expression of Jagged 2. Specifically, it refers to compositions useful for inhibiting the expression of Jagged 2, a human homologue of the Drosophila Scrate gene, which is involved in differentiation and cell fate, as well as positive feedback control over signalling genes such as Notch 1, Notch 3 and Jagged 1. The Jagged 2 gene is located on chromosome 14932, a region that has been implicated in genetic diseases including Usher syndrome type la that is associated with cetinitis pigmentosa. The present invention describes antisense oligonucleotides that comprise at least one modified sugar moiety, a 2'-O-methoxyethyl (2' MOE) and at least one modified nucleobase, a 5-cmethoxyethyl (2' MOE) and at least one modified nucleobase, a 5-cmethoxyethyl (2' MOE) and at least one modified nucleobase, a 5-cmethylcytosine. These compounds are useful for modulating the mucleic acid molecules encoding Jagged 2, ultimately modulating the amount of Jagged 2 produced, which in turn is useful for research
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New antisense oligonucleotides of 8-40 nucleobases, useful for modulating the function of nucleic acid molecules encoding Jagged 2, ultimately modulating the amount of Jagged 2 produced.
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                                                                                                                                                                                                                                                                                                                                                                                       human, Jagged 2; differentiation; cell fate; signalling;
Usher syndrome type la; retinitis pigmentosa; PCR; ss; primer.
                    3.2%; Score 14.4; DB 1; Length 16; 93.8%; Pred. No. 2.8e+02; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         primer used to amplify human Jagged 2 DNA of the invention
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                                                                                                                                                                                                                              ADH57064 standard; DNA; 16 BP
                                                                                                                      1 CCCAGGGCTTCTCCGG 16
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                                                                                                  265 CCCGGGGCTTCTCCGG
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                                                            Conservative
Query Match
Best Local Similarity
Local 15; Conserve
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (FREI/) FREIER S M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US2003207839-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
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                                                                                                                                                                                              RESULT 321
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Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    prostate, colorectal, brain, oesophageal, stomach, bladder, pancreatic, cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or multidrug resistant cancer. The method involves use of cother drug therapies such as monoclonal antibodies, REL-A-specific inhibitors or chemotherapy including paclitaxel, docetaxel, cisplatin, methotrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edatrexate, gemcitabine or radiation therapy. The enzymatic and antisense nucleic acid molecules are also useful for treating inflammatory disease such as rheumatoid arthritis, restenosis, asthma, Crohn's disease, diabetes, obesity, autoimmune disease, lupus, multiple sclerosis, transplant/graft rejection, gene therapy applications, ischaemia/reperfusion injury (central nervous system (CNS) and myocardial), glomerulonephritis, sepsis, allergic airway inflammatory bowel disease or injury application, inflammatory bowel disease or included the contral nervous systems of the contral nervous systems.
                                                                                                                                                                                           G-cleaver; amberzyme; cancer; REL-A activity; breast cancer; human; lung cancer; prostate cancer; colorectal cancer; brain cancer; coesphageal cancer; scancer; colorectal cancer; pancreatic cancer; cervical cancer; head and neck cancer; ovarian cancer; melanoma; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; chemotherapy; paclitaxel; docetaxel; cisplatin; methotrexate; gencitabine; radiation therapy; inflammatory disease; asthma; diabetes; theumatoid arthritis; restenosis; Crohn's disease; obesity; ischaemis; gene therapy; autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/graft rejection; reperfusion injury; glomerulonephritis; allergic airway inflammatory bowel disease; infection; se.
                                                                                                                                                                         Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             infection. This sequence represents the substrate of a novel enzymatic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Draper KG;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             92US-00987132.
94US-00245466.
94US-00291932.
96US-00777916.
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MCSWIGGEN J.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              JS2002177568-A1
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15-AUG-1994;
23-DEC-1996;
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Gaps

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3.2%; Score 14.4; DB 1; Length 16; 93.8%; Pred. No. 2.8e+02; ve 0; Mismatches 1; Indels

llarity 93.8%; Conservative 0

Best Local Similarity Matches 15; Conserv

Query Match

265 CCCGGGGCTTCTCCGG 280

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CCCAGGCTTCTCCGG 16

RESULT 322

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03-JUN-2004
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(MACE/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                         Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme; G-cleaver; amberzyme; cancer; REL-A activity; breast cancer; human; lung cancer; prostate cancer; colorectal cancer; brain cancer; coesophageal cancer; stomach cancer; bladder cancer; pancreatic cancer; cervical cancer; had and neck cancer; ovarian cancer; melanoma; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; common partial cancer; pactitates; pactitates; gencitabine; radiation therapy; inflammatory disease; asthma; diabetes; gencitabine; radiation therapy; inflammatory disease; asthma; diabetes; rheumatorid arthritis; restenosis; Crohn's disease; obesity; ischaemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/graft rejection; reperfusion injury; glomerulonephritis; allergic airway inflammation; inflammatory bowel disease; infection; ss.
                                                                                                             Gaps
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                                                                      3.2%; Score 14.4; DB 1; Length 17; 75.0%; Pred. No. 3e+02; tive 3; Mismatches 1; Indels
                                     Sequence 17 BP; 0 A; 10 C; 4 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                             NFKB sub-unit modulating inozyme substrate #145.
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94US-00291932.
96US-00777916.
                                                                                                                                                    131 CCTCGGCCTGCCGCCT 146
                                                                                                                                                                                                                                                                                    ACA06326 standard; RNA; 17 BP
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                                                                                                               12; Conservative
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MCSWIGGEN J.
DRAPER K G.
   nucleic acid molecule
                                                                                             Best Local Similarity
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23-DEC-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-NOV-2002
                                                                                                                                                                                                                                                                                                                         ACA06326;
                                                                          Query Match
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                                                                                                                  Matches
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prostate, colorectal, brain, oesophageal, stomach, bladder, pancreatic, cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or multidrug resistant cancer. The method involves use of other drug therapies such as monoclonal antibodies, REL-A-specific inhibitors or chemotherapy including paclitaxel, docetaxel, cisplatin, methotrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, methotrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, methotrexate, committabine or radiation therapy. The enzymatic and antisense nucleic acid molecules are also useful for treating inflammatory disease such as the molecules are also useful for treating inflammatory disease such as checkin, general methods, restenosis, statema, Crohn's disease, diabetes, obesity, autoimmune disease, lugus, multiple sclarosis, transplant/graft rejection, gene therapy applications, ischaemia/reperfusion injury sepsis, allergic airway inflammatory bowel disease or infection. This sequence represents the substrate of a novel enzymatic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ss; enzymatic nucleic acid; RNA cleavage; hepatitis C virus; HCV;
HCV infection; type I interferon; DNAzyme.
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Pred. No. 3e+02;
3; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 17 BP; 0 A; 11 C; 3 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pavco PA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; SEQ ID NO 800; 198pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3;
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Best Local Similarity 75.0°
Matches 12, Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              nucleic acid molecule
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hepatitis C virus.
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24-FEB-2003 (first entry)
                                                                                                                                                                                                               Unidentified,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ig molety
                    AAD47637;
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present sequence is an antisense oligonucleotide which was used in an RNase mapping experiment. This enables the identification of sites within the Ii RNA strand which hybridise to antisense DNA. These sites can then be used as targets for antisense strands which may, using gene therapy, be used as tumour cell vaccines (for example to treat carcinomas, namelanoma, laukaemia, lymphomas, stomach, breast, colon or rectum, lung, prostate, bladder, pancreas, brain and ovarian cancers), or they can be used to treat autoimmune diseases including rheumatoid arthritis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Immunoregulator; antisense oligonucleotide; cancer; tumour cell vaccine; rheumatoid arthritis; autoimmune disease; diabetes mellitus; thyroiditis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Antisense oligonucleotide for use in RNase H mapping assay SEQ ID NO: 24.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cancer cell vaccine for treating malignancies, autoimmune disorders and isolating autodeterminant peptides comprises a regulator of invariant chain protein expression or immunoregulatory function.
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                                                                                Query Match 3.2%; Score 14.4; DB 1; Length 17; Best Local Similarity 82.4%; Pred. No. 3e+02; Matches 14; Conservative 1; Mismatches 2; Indels
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                                          Sequence 17 BP; 5 A; 6 C; 4 G; 0 T; 1 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 1; Page 46; 94pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      diabetes mellitus and thyroiditis
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                                                                                                                                                                                                             1 AGGACUNGGCCCACACA 17
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Best Local Similarity
Local Siconservative
Local 15; Conservative
                                                                                                                                                                                                                                                                                                                        AAA63120 standard; DNA; 18
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AAD47637/c
ID AAD476
                                                                                                                                                                                                                                                                                RESULT 325
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                                                                                   Immunoglobulin; Ig; antibody; immunocytokine; immunofusion; immunoligand;
protease resistance; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       we fusion protein, useful for producing antibodies, immunocytokines or tusion proteins that enhance the expression or protesse resistance of fusion protein, comprises an immunoglobulin (Ig) moiety fused to a non-
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intron; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention relates to methods and compositions for efficiently
Forward primer, to construct human KSgamma2h antibody expression vector.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 5; Page 63; 71pp; English.
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nes 15; Conserv
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The present invention relates to detecting and identifying fungal pathogenic species in a sample. The method involves hybridizing a nucleic acid of a fungal pathogen possibly present in the sample with at least one oligonucleotide probe, from an Internal Transcribed Spacer (ITS) region. The method is useful for simultaneous detection and differentiation of clinically important fungi in a single assay, particularly Candida albicans, C. parapsilosis, C. tropicalis, C. kefyr, C. krused, C. glabrata, C. dubliniensis, Aspergillus flavus, A. versicole, A. nidulans, A. fumigatus, C. neoformans and pneumocystis carini. The method is especially useful in the detection of opportunistic infections in patients with impaired immunity systems, such as organ transplants patients, patients receiving intensive anticancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The patent discloses new macromolecules of formula PNA-DNA-PNA, in which DNA comprises at least one 2'-deoxynucleotide and each PNA comprises at least one peptide nucleic acid subunit. These compounds have increased resistance to nuclease and increased specific binding affinity, and they can activate RNaseH for target strand cleavage. They can hybridise specifically to a nucleic acid strand (especially RNA) and are useful (1) for treating diseases associated with undesirable production of protein,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New chimeric macromolecules contg. DNA and peptide nucleic acid segments - with good nuclease stability and binding affinity, also activating RNaseH, useful for treating disease, in diagnosis and for identifying
ouggood and Aspergillus, comprises hybridizing the amplified nuclacid of the fungal pathogen with a probe from the internal transcribed spacer region of a DNA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Oligonucleotide used in PNA-DNA-PNA chimeric macromolecule.
                                                                                                                                                                                                                                                                                                                                                                     DB 1; Length 17; 3.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                   0; Indels
                                                                                                                                                                                                                                                                                                                                      Sequence 17 BP; 1 A; 7 C; 7 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                     3.1%; Score 14; DB 100.0%; Pred. No. 3.3 tive 0; Mismatches
                                                                                                                                                                                                                                                                                                          treatments, diabetics or AIDS patients
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                                                                             Claim 1; Page 46; 59pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                     328 CTCTCGGGGGGGAG 341
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAV81598 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2 crcrceeeeeceae 15
                                                                                                                                                                                                                                                                                                                                                                      Query Match 3.1
Best Local Similarity 100.
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         chemotherapeutic agents.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1995-206893/27.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  11-MAY-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAV81598;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cook PD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 329
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAV81598
                g
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
                                                                                                                                                                                                                                                                                                                                     The invention relates to a method of identifying a modulator of Cdc25 activity that comprises contacting a test cell having a recombinant Cdc25 phosphatase gene whose expression alters transcription of a selected gene, with a compound under conditions where recombinant Cdc25 phosphatase gene is expressed and alters the transcription of a selected gene as an indication of the compound being a modulator of Cdc25-mediated transcription. The method is useful for identifying modulators of Cdc25
                                                                                                                                                                                                                                        Assay for screening modulators of Cdc25 activity by using a cell having recombinant Cdc25 phosphatase gene whose expression alters the transcription of a selected gene in the presence of a modulator.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     activity. Sequences AAF57363-376 represent intron/exon splice junction sequences of the murine Cdc25A gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Van Der Weide M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Detecting and identifying fungal pathogens, especially Candida
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 14; DB 1; Length 17;
Pred. No. 3.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             pathogenic; Internal Transcribed Spacer; ITS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Rossau R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Fungal pathogenic species identification probe #21.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 17 BP; 4 A; 3 C; 4 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (INNO-) INNOGENETICS NV.
(IRBI-) ENTERPRISE IRELAND T/A BIORESEARCH IRELA.
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                                                                                                                                                                                                                                                                                                           Example 1; Page 15; 55pp; English
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                                                                                                                  99US-0153639P
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                                                                                  11-SEP-2000; 2000WO-US024838
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        155 TCATTCTAGAGCAA 168
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                                                                                                                                                                                Timm J;
                                                                                                                                                (BADI ) BASF AG
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                    WO200120034-A2
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11-JUN-1999;
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                                                    22-MAR-2001
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AAX6295

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New nucleic acid(s) for regulating the Grb2-related with Insert Domain (GRID) gene comprises using antisense and enzymatic nucleic acid
                                                                                                                                                                                                                                                                                                                   Human, Grb2-related with Insert Domain; GRID; T-cell;
co-etimulatory adaptor protein; tissue rejection; graft rejection;
leukaemia; cytostatic; ss.
                                                                                                                                                                                                                                                                                      Human GRID NCH ribozyme substrate oligonucleotide #151
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mcswiggen JA,
Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              molecules such as hammerhead ribozymes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 4; Page 65; 108pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           used to illustrate the invention
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                                      106 CGCTGACTTTCAGCGGG 122
                                                                                                                                                                        BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3.1%;
88.2%;
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                                                                          cecueccurucaecues
                                                                                                                                                                      ABL46697 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (RIBO-) RIBOZYME PHARM INC. (GLAX ) GLAXO GROUP LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 200 CCTCCCGGGGACCTGCG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABL46496 standard; RNA; 17
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                                                                                                                                                                                                                                                    (first entry)
  Conservative
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Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200162911-A2
                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    27-JUN-2003
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  11;
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                                                                                                                                                                                                              ABL46697;
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                                                                                                                                    RESULT 331
                                                                                                                                                      ABL46697/C
  Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention describes an enzymatic nucleic acid molecule (I) with RNA cleaving activity, which modulates the expression of a plant some. Also described is a gene comprising a cDNA sequence encoding maize Delta-9 desaturase. (I) can be used to modulate expression of a gene, preferably Delta-9 desaturase or a granule bound starch synthase (GBSS) gene, in a plant (preferably a maize or canola plant). (I) can be used to modulate caffeine synthesis in a coffee plant, nicotine production in a tobacco plant, fruit ripening processes in an apple, tomato, pear, plum or peach plant, flower pigmentation in a rose, petunia, chrysanthemum or marigold plant or lignin production in a tobacco, aspen, poplar or pine
                    for
                  contacting a test solution with the macromolecule and RNaseH), or (3) for in-vivo enhancement of polynucleotide hybridisation and RNase activity. They can also be used diagnostically and for screening chemotherapeutic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Maize; corn; Zea mays; delta-9 desaturase; GBSS; target; substrate; granule bound starch synthase; hammerhead ribozyme; hairpin ribozyme; modulation; gene expression; transgenic plant; cleavage; canola plant; caffeine synthesis; coffee plant; nicotine production; tobacco; fruit ripening; flower pigmentation; lignin production; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Skokut TA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ribozyme which modulates plant gene expression - preferably modulates expression of DELTA-9 desaturase or granule bound starch synthase in maize or canola.
                                                                                                                                                                                            Gaps
  in-vitro modification of sequence-specific nucleic acid (by
                                                                                                                                                                                            ö
                                                                                                                                                    3.1%; Score 13.8; DB 1; Length 17; 88.2%; Pred. No. 3.4e+02; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 13.8; DB 1; Length 17; Pred. No. 3.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Delta-9 desaturase hamerhead ribozyme target SEQ ID NO:826.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Merlo PAO,
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                                                                                                                Sequence 17 BP; 2 A; 5 C; 3 G; 7 T; 0 U; 0 Other;
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Merlo DJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 38; Page 86; 155pp; English.
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                                                                                                                                                                                                                                                                                                                                                                     AAX62951 standard; RNA; 17
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                                                                                                                                                                      Local Similarity 88.2
nes 15; Conservative
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Folkerts O,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1997-202224/18
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Zea mays.
                                                                                                                                                                                                                                  102
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                                                                                                                                                      Query Match
                                                                              agents
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Ellis JH;

Hamblin PA,

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The present invention relates to oligonucleotides that downregulate the expression of human Grb2-related with Insert Domain (GRID) gene. GRID is a T-cell co-stimulatory adaptor protein. The oligonucleotides are useful for modulating the expression of GRID, to treat conditions such as tissue/graft rejection and leukaemia. The oligonucleotides can also be administered in conjunction with other therapies such as radiation, chemotherapy and cyclosporin treatment. The present oligonucleotide was
                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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Pred. No. 3.4e+02;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                  Seguence 17 BP; 3 A; 4 C; 9 G; 0 T; 1 U; 0 Other;
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ABL46496/C
ID ABL4649
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AC ABL464
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DT 27-JUN
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                                                                                                                                                                                                                                                                                                                                                      The present invention relates to oligonucleotides that downregulate the expression of human Grb2-related with Insert Domain (GRLD) gene. GRLD is a T-cell co-stimulatory adaptor protein. The oligonucleotides are useful for modulating the expression of GRLD, to treat conditions such as tissue/graft rejection and leukaemia. The oligonucleotides can also be administered in conjunction with other therapies such as radiation, chemotherapy and cyclosporin treatment. The present oligonucleotide was used to illustrate the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; hammerhead ribozyme; cytostatic; antitumour; antidiabetic; ophthalmological; antiarthritic; antipsoriatic; virucide; osteopathic; vulnerary; cancer; lymphoma; Ewing's sarcoma; melanoma; psoriasis; tumour angiogenesis; diabetic retinopathy; macular degeneration; neovascular glaucoma; myopic degeneration; arthritis; verruca vulgaris; angiofibroma of tuberous solerosis; port wine stain; wound healing; Sturge Weber syndrome; Kippel-Trenaunay-Weber syndrome; leukaemia; ss; osler-Weber-rendu syndrome, leukaemia; osteoporosis; DNAzyme; inozyme;
                                                                                                                                                                                                                                                                                 Domain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
       Human, Grb2-related with Insert Domain, GRID; T-cell; co-stimulatory adaptor protein; tissue rejection; graft rejection; leukaemia; cytostatic; ss.
                                                                                                                                                                                                                              Ellis JH
                                                                                                                                                                                                                                                                              New nucleic acid(s) for regulating the Grb2-related with Insert E (GRID) gene comprises using antisense and enzymatic nucleic acid molecules such as hammerhead ribozymes.
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                                                                                                                                                                                                                              Mcswiggen JA, Hamblin PA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human ERG Amberzyme target sequence Seq ID No 1949.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 17 BP; 3 A; 5 C; 8 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                  Claim 4; Page 61; 108pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   201 CTCCCGGGGACCTGCGG 217
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                                                                                                                                       23-FEB-2001; 2001WO-US005957
                                                                                                                                                                                                                             Jarvis T, Von Carlowitz I,
                                                                                                                                                                                      (RIBO-) RIBOZYME PHARM INC.
(GLAX ) GLAXO GROUP LTD.
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hes 15; Conservative
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                                                                                      WO200162911-A2.
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                                                              Homo sapiens
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ABK19302/c
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The invention relates to a nucleic acid molecule (I) which down regulates expression of an Ets-related gene (ERG). (I) is useful for treating conditions selected from cancer, lymphoma, Ewing's sarcoma, melanoma, tumour anglogenesis, diabetic retinopathy, macular degeneration, tumour anglogenesis, diabetic retinopathy, macular degeneration, cumour anglofibroma of tuberous sclerosis, port-wine stains, Sturge wolfares, anglofibroma of tuberous sclerosis, port-wine stains, Sturge Weber syndrome, Rippel-Trenaunay-Weber syndrome, Osler-Weber-rendu (Syndrome, leukaemia, osceoporosis and wound healing. (I) is useful for treating a patient having a condition associated with the level of ERG, by contacting cells of the patient with (I) under conditions suitable for the treatment. Leukaemia or tumour angiogenesis is treated by administering (I) to the patient in conjunction with one or more of other therapies such as radiation or conjunction with one or more of other therapies such as radiation or chemotherapy treatment. (I) is useful for reducing ERG activity in a cell, by contacting the cell with (I). (I) is useful for cleaving RNA of ERG gene, by contacting (I) with RNA, in the presence of a divalent cation such as MG2+. (I) is useful for diagnosis of conditions and diseases related to the expression of ERG, and as diagnostic tool to examine genetic drift and mutations within diseased cells or to detect the presence of ERG RNA in a cell. (I) is useful for specifically targeting genes that share homology with ERG gene or ERG fusion genes. ARX.7354-ABXZ2719 represent mucleic acide, including antisense and enzymatic mucleic acide molecules which regulate expression of ERG, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ1; MDZ12; chromosome 7g22.1; chromosome 6p21.3-22.2; chromosome 6p21.3-22.3; chromosome 16p11.2; chromosome 15g26.1; cancer; developmental disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                  Novel polynucleotide which down regulates expression of Ets-related gen useful for treating cancer, diabetic retinopathy, macular degeneration, arthritis, psoriasis, verruca vulgaris and Sturge Weber syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                      Randi AM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ;
0
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                                                                                                                                                                                                                                      Von Carlowitz I, Mcswiggen JA, Mclaughlin F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 17 BP; 1 A; 5 C; 6 G; 0 T; 5 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 13.8; DB 1;
Pred. No. 3.4e+02;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human MDZ3 scanning oligonucleotide SEQ ID 814.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       related PCR primers of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 4; Page 125; 149pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 140
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                                                               16-MAY-2000; 2000US-00572021.
16-MAY-2001; 2001WO-US015866.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Buet Local Similarity 88.20,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 124 GGAAAAGCCTCGGCCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17 GGAAAAGCCTCCGCCAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                            (RIBO-) RIBOZYME PHARM : (GLAX ) GLAXO GROUP LTD
                                                                                                                                                                                                                                                                                                  WPI; 2002-082995/11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EP1281758-A2.
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                                                                                                                                                                                                                                      Jarvis T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADA99825;
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                                                                                                                                                                                                                                                                                                      proteins and their coding sequences: MD23, MD24, MD27, MD212. MD23 be cancoded at chromosome 792.1, MD24 is encoded at chromosome 692.1, MD24 is encoded at chromosome 692.13.22.2, MD24.1 is encoded at chromosome 692.13.22.2, MD24 is encoded at chromosome 692.13.22.2, MD24.1 is encoded at chromosome 1591.3 encoded at chromosome 1592.6.1. The MD23, MD24, MD27, and MD21 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MD23, MD24, MD27, or MD212, e.g. cancer or developmental disorders. The nucleic caused by altered expression of MD23, MD24, MD27, or MD212. The nucleic acids can also be used as probes to detect and characterize gross alterations in MD23, MD27, or MD212 genetic locus. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nucleic acid molecule; Hepatitis C virus; HCV; Hepatitis B virus; HBV; RNA stability; RNA expression; RNA synthesis; antisense; enzymetic nucleic acid; hammerthead ribozyme; DNAzyme; inozyme; inozyme; amberzyme; G-cleaver ribozyme; decoy molecule; aptamer; HBV reverse transcriptase; Enhancer I region; viral replication; degenerative; disease state, HBV infection; HCV infection; cirrhosis; liver failure; hepatocellular carcinoma; hepatotropic; cytostatic; virucide; antiinflammatory; substrate; ss.
                                                                                                                                                                                      New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                           The present invention relates to novel human zinc finger-containing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3.1%; Score 13.8; DB 1; Length 17; 88.2%; Pred. No. 3.4e+02; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 17 BP; 2 A; 2 C; 8 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                               Example 8; SEQ ID NO 814; 103pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HCV DNAzyme substrate sequence #800.
                                                                                                                                                                                                                                  MDZ4, MDZ7 or MDZ12, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      24 AGGGGTGGTGGCCATTT 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AGGGGTTGGGGCCATTT 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ACD58382 standard; RNA; 17 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  26-MAR-2002; 2002WO-US009187.
                                         30-JUL-2002; 2002EP-00016874
                                                                      02-AUG-2001; 2001US-00922181
                                                                                                                               Shannon M, Gu Y, Nguyen C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15; Conservative
                                                                                                                                                            WPI; 2003-423107/40.
                                                                                                   (AEOM-) AEOMICA INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hepatitis C virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200281494-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   24-SEP-2003
             05-FEB-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17-0CT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ACD58382;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 335
8
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the synthesis, expression and/or grability of Hepatitis C virus (HCV) or Hepatitis B virus (HBV) RNA. The nucleic acid molecules include antisense and enzymatic nucleic acids such as hammerhead ribozymes, DNAzymes, incorpuse, zinzymes, amberzymes, and G-cleaver ribozymes, DNAzymes, controlled acids such an aptamers that bind to HBV reverse transcriptase primer sequences, as well as oligonucleotides that specifically bind the Enhancer I region of HBV DNA. The nucleic acids may be used to modulate the expression of HBV genes and HBV viral replication. Also disclosed is a method for screening compounds and/or potential therapies directed against HBV, and compounds that modulate the expression and/or replication of HCV. The compounds and control are useful for the treatment of degenerative and disease states related to HBV and HCV infection, replication and gene carcinoma. The present sequence represente a substrate for one of the HCV DNAzyme or minus strand DNAzyme sequences disclosed in the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nucleic acid molecule, Hepatitis C virus, HCV; Hepatitis B virus; HBV; RNA stability; RNA expression; RNA synthesis, antisense; enzymatic nucleic acid; hammerhead ribozyme; DNAzyme; inozyme; zinzyme; amberzyme; G-cleaver ribozyme; decoy molecule; aptemer; HBV reverse transcriptase; Enhancer I region; viral replication;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to nucleic acid molecules which modulate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 Lee
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3.1%; Score 13.8; DB 1; Length 17; 82.4%; Pred. No. 3.4e+02; ive 1; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pavco P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HCV minus strand DNAzyme substrate sequence #853.
                                                                                                                                                                                                                                                                                                                                                                                                                                           Mcswiggen J, Morrissey D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 17 BP; 5 A; 6 C; 5 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1; Page 248; 387pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              432 AGGACTCGGCTCACACA 448
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AGGACUGGCCCCACACA 17
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26-MAR-2001; 2001US-00817879.
08-JUN-2001; 2001US-0087478.
28-JUN-2001; 2001US-0296876P.
24-OCT-2001; 2001US-0337055P.
05-DEC-2001; 2001US-0337055P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity 82.4%;
Les 14; Conservative
                                                                                                                                                                  RIBOZYME PHARM INC.
BLATT L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Roberts E;
                                                                                                                                                                                                                  MACEJAK D.
MCSWIGGEN J.
MORRISSEY D.
PAVCO P.
                                                                                                                                                                                                                                                                                                                                                                                                                                              Macejak D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2003-229207/22
                                                                                                                                                                                                                                                                                                                                                        DRAPER K.
ROBERTS E.
                                                                                                                                                                                                                                                                                                                                    LEE P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24-SEP-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Blatt L,
                                                                                                                                                                                                                                                                                                                               (LEEP/)
(DRAP/)
(ROBE/)
                                                                                                                                                                  (RIBO-)
                                                                                                                                                                                              (BLAT/)
(MACE/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Draper
                                                                                                                                                                                                                                                       (MCSM/
                                                                                                                                                                                                                                                                              (MORR/
                                                                                                                                                                                                                                                                                                              PAVC/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ACD63046/C
ID ACD630
AC ACD630
XX
XX
DT 24-SEP
XXX
DE HCV mi
XX
KW Nuclei
KW RNA st
KW enzyme
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Matches
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Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; murine; tumour suppression; tumour reversion; apoptosis; virus resistance; viral disease; tumour; cell degeneration; cancer; Alzheimer's disease;

schizophrenia; ss.

WO2003025176-A2. Mus musculus,

27-MAR-2003

Murine oligonucleotide associated with tumour supression, SEQ ID 5862.

01-JUL-2003 (first entry)

ACC68615;

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Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus
degenerative; disease state; HBV infection; HCV infection; cirrhosis;
         liver failure; hepatocellular carcinoma; hepatotropic; cytostatic; virucide; antiinflammatory; substrate; ss.
                                                                                                                                                                                                                                                     Pavco P, Lee P;
                                                                                                                                                                                                                                                   Mcswiggen J, Morrissey D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 17 BP; 3 A; 11 C; 2 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                              Claim 1; Page 290; 387pp; English.
                                                                                                       26-MAR-2001; 2001US-00817879.
08-UUN-2001; 2001US-00877478.
08-UUN-2001; 2001US-0296876P.
24-0CT-2001; 2001US-0335059P.
05-DEC-2001; 2001US-0337055P.
                                                                                        26-MAR-2002; 2002WO-US009187
                                                                                                                                                           RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                  Macejak D,
Roberts E;
                                                                                                                                                                                     MCSWIGGEN J.
MORRISSEY D.
                                                                                                                                                                                                                                                                            WPI; 2003-229207/22.
                                                                                                                                                                             MACEJAK D.
                                                                                                                                                                                                      PAVCO P.
LEE P.
DRAPER K.
ROBERTS E.
                                   Hepatitis C virus.
                                                     WO200281494-A1
                                                                                                                                                                                                                                                          Draper K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    invention
                                                                                                                                                                    (BLAT/)
(MACE/)
(MCSW/)
(MORR/)
(PAVC/)
                                                                                                                                                                                                                        (DRAP/)
(ROBE/)
                                                                                                                                                                                                                                                  Blatt L,
                                                                                                                                                           RIBO-)
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The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HCV) or Hepatitis B virus (HBV) RNA. The nucleic acid molecules include antisense and enzymatic nucleic acids such as hammerhead ribozymes, DNAzymes, or inozymes, zinzymes, amberzymes, anderzymes, anderziptase and or-leaver ribozymes. Also disclosed are nucleic acids may be used to modulate the expression of HBV CC and of Compounds and HBV viral replication, Also disclosed is a method for screening compounds and/or potential therapies directed against HBV, and compounds that modulate the expression and/or replication of HCV. The compounds and disease states related to HBV and HCV infection, replication and gene expression such as cirrhosis, liver failure, and hepatocallular expression such as cirrhosis, liver failure, and hepatocallular expression and number sequence represents a substrate for one of the HCV invention in the present
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The present invention relates to murine oligonucleotides (ACC62754-ACC68006), which are associated with tumour suppression, tumour reversion, apoptosis and virus resistance. The oligonucleotides are useful as (1) as probes and primers for detecting, identifying, quantifying and/or amplifying nucleic acid, e.g. as one component of a gene chip; in vitro as (anti) sense reagents; and (2) for production of recombinant polypeptides. The oligonucleotides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell degeneration, specifically cancer but also Alzheimer's disease and schizophrenia

New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies

Tuijnder

Amson R,

Telerman A,

WPI; 2003-333167/31.

(MOLE-) MOLECULAR ENGINES LAB.

17-SEP-2002; 2002WO-IB004210. 17-SEP-2001; 2001FR-00011979. Disclosure; Page 716; 738pp; French.

with tumors and cell dand transfected cells.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 antisense oligonucleotide; neurite growth inhibitor; NOGO; prostagalandin D2 receptor; PTGDR; IkappaB kinase; IKK; protein kinase PKR; cerebrovascular accident; protein kinase PKR; cerebrovascular accident; central nervous system injury; CNS injury; spinal cord injury; cancer; melanoma; lymphoma; glioma; inflammatory disease; rheumatoid arthritis; restenciss; asthma; Crohn's disease; diabetes; obesity; autoimmune disease; lupus; multiple sclerosis; transplant rejection; graft rejection; schaemia; reperfusion; glomerulonephritis; sepsis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3.1%; Score 13.8; DB 1; Length 17;
88.2%; Pred. No. 3.4e+02;
ve 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human NOGO receptor zinzyme substrate sequence #84.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 17 BP; 1 A; 6 C; 2 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                368 AGGAAGAGGAACGGAGC 384
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADL47097 standard; RNA; 17 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 88.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17 AGGAAAGGAACGGATC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity 88.2
nes 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20-MAY-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 338
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Gaps

; 0

3.1%; Score 13.8; DB 1; Length 17; 38.2%; Pred. No. 3.4e+02; Ive 0; Mismatches 2; Indels

88.2%;

Conservative

Local Similarity les 15; Conserv

Matches

Query Match

TGGGAGGGGTGGTGGCC 36

20

17 redeAdderederedece 1

g

BP.

ACC68615 standard; DNA; 17

RESULT 337

ACC68615/ ID ACC6

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asthma; allergic rhinitis; atopic dermatitis; human PTGDR;
                                                                Unidentified
  allergy; as
substrate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADL48304;
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that down regulate the expression or inhibit the function of a receptor
for a neurite growth inhibitor, NOGO, prostaglandin D2 receptor (PTGDR),
IKappaB kinase (IKK), or protein kinase PKR. The nucleic acids of the
invention are useful for treating: cerebrovascular accident, central
nervous system (CNS) injury, spinal cord injury, cancer (e.g. melanoma,
lymphoma or glioma), inflammatory disease (e.g. rheumatoid arthritis,
restenosis or asthma), inflammatory disease (e.g. rheumatoid arthritis,
cestenosis or asthma), inflammatory disease, diabetes, obesity, autoimmune
ischaemia/reperfusion injury, glomerulonephritis, sepsis, and allergic
conditions (e.g. asthma, allergic rhinitis or atopic dermatitis). The
nucleic acids of the invention are also useful for down-regulating the
cexpression of a target gene and as a diagnostic tool to examine genetic
drifts and mutations within diseased cells or to detect the presence of a
target RNA in a cell. The present RNA sequence represents a human NOGO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel enzymatic nucleic acid that down-regulates expression of neurite growth inhibitor receptor, prostaglandin D2 receptor, IkappaB kinase or protein kinase PKR genes, for treating cancer and inflammatory disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                antisense oligonucleotide; neurite growth inhibitor; NOGO; prostaglandin D2 receptor; PTGDR; IkappaB kinase; IKK; protein kinase PKR; cerebrovascular accident; central nervous system injury; CNS injury; spinal cord injury; cancer; melanoma; lymphoma; glioma; inflammatory disease; rheumatoid arthritis; restenosis; asthma; Crohn's disease; diabetes; obesity; autoimmune disease; lupus; multiple sclerosis; transplant rejection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       graft rejection; ischaemia; reperfusion; glomerulonephritis; sepsis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                                                                                                                  Haeberli P, Mcswiggen J, Fosnaugh K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ch 3.1%; Score 13.8; DB 1; Length 17; 11 Similarity 76.5%; Pred. No. 3.46+02; 13; Conservative 2; Mismatches 2; Indels
allergy, asthma, allergic rhinitis, atopic dermatitis, NOGO receptor zinzyme; substrate; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 17 BP; 0 A; 6 C; 8 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 9; SEQ ID NO 630; 317pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human PTGDR substrate sequence #1040.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      receptor zinzyme substrate sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 263 GGCCCGGGGCTTCTCCG 279
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                                                                                                                                                                                                                                 05-APR-2001; 2001US-00827395.
29-MAY-2001; 2001US-0294412P.
28-AUG-2001; 2001US-0315315P.
                                                                                                                                                                                            03-APR-2002; 2002WO-US010512
                                                                                                                                                                                                                                                                                                                        (RIBO-) RIBOZYME PHARM INC.
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                                                                                                                                                                                                                                                                                                                                                                Blatt L, Chowrira B,
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Best Local Similarity
                                                                                                          WO200281628-A2
                                                                Unidentified
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ID ADL51
XX
AC ADL51
XX
XY
DT 20-M
XX
KW antick
KW pros
KW pros
KW cent
KW mela
KW mela
KW rest
KW auto
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The invention comprises nucleic acids (e.g. antisense oligonucleotides)
that down regulate the expression or inhibit the function of a receptor
for a neurite growth inhibitor, NOGO, prostaglandin DZ receptor (FTGDR),
IKappa Kinase (IKK), or protein Kinase PKR. The nucleic acids of the
invention are useful for treating: cerebrovascular accident, central
nervous system (CNS) injury, spinal cord injury, cancer (e.g. melanoma,
nervous system (CNS) injury, spinal cord injury, cancer (e.g. melanoma,
restenosis or asthma), Crohn's disease, diabetes, obesity, autoimmune
disease, lupus, multiple sclerosis, transplant/graft rejection,
cischaemia/reperfusion injury, glomerulonephritis, sepsis, and allergic
conditions (e.g. asthma, allergic rhinitis or atopic dermatitis). The
nucleic acids of the invention are also useful for down-regulating the
cxpression of a target gene and as a diagnostic tool to examine genetic
drifts and mutations within diseased calls or to detect the presence of a
target RNA in a cell. The present RNA sequence represents a human PKR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      antisense oligonucleotide; neurite growth inhibitor; NOGO; prostaglandin D2 receptor; PTGDR; IkappaB kinase; IKK; prostaglandin D2 receptor; PTGDR; IkappaB kinase; IKK; protein kinase PKR; cerebrovascular accident; central nervous system injury; CNS injury; spinal cord injury; cancer; melanoma; lymphoma; glioma; inflammatory disease; rheumatoid arthritis; restenoisis; asthma; Crohn's disease; diabetes; obesity; autoimmune disease; lupus; multiple sclerosis; transplant rejection; graft rejection; ischaemia; reperfusion; glomerulonephritis; sepsis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel enzymatic nucleic acid that down-regulates expression of neurite growth inhibitor receptor, prostaglandin D2 receptor, IkappaB kinase or protein kinase PKR genes, for treating cancer and inflammatory disease.
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                                                                                                                                                                                                                                                                                                                                                                                                      Fosnaugh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Seguence 17 BP; 9 A; 2 C; 5 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                  Haeberli P, Mcswiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 161; SEQ ID NO 5454; 317pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human IKK-gamma substrate sequence #814.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CTGTTTTTCTCGCTGAC 112
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                                                                                                                                                                                          05-APR-2001; 2001US-00827395.
29-MAY-2001; 2001US-0294412P.
28-AUG-2001; 2001US-0315315P.
                                                                                                                                  03-APR-2002; 2002WO-US010512
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Best Local Similarity 88.20,
Best Local Similarity 18.20,
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADL48304 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                   (RIBO-) RIBOZYME PHARM INC.
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                                                                                                                                                                                                                                                                                                                                                                                                      Blatt L, Chowrira B,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                substrate sequence.
WO200281628-A2
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Page 174

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allergy; asthma; allergic rhinitis; atopic dermatitis; Human IKK-gamma;
                                                                                                                                                                            05-APR-2001; 2001US-00827395.
29-MAY-2001; 2001US-0294412P.
28-AUG-2001; 2001US-0315315P.
                                                                                                                                                  03-APR-2002; 2002WO-US010512.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 GCGUGGAGGCCGCCCC 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 76.5%;
Matches 13; Conservative
                                                                                                                                                                                                                                         (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   246 GCCTGGAGGCCGCGGTC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             gamma substrate sequence.
                                                                                                                                                                                                                                                                          Chowrira B,
                                                                                                                                                                                                                                                                                                     WPI; 2003-058513/05.
                                                                                     WO200281628-A2.
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                           substrate; ds.
                                                        Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                    17-0CT-2002,
                                                                                                                                                                                                                                                                        Blatt L,
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that down regulate the expression or inhibit the function of a receptor
for a neurite growth inhibitor, NOGO, prostaglandin D2 receptor (PTGDR),
[KappaB kinase (IKK), or protein kinase PKR. The nucleic acids of the
invention are useful for treating: creebrovascular accident, central
nervous system (CKS) injury, spinal cord injury, cancer (e.g. melanoma,
[KappaB kinase (IKK)], inflammatory disease (e.g. rheumatoid arthritis,
restencis or asthma), Crohn's disease, dispetes, obesity, autoimmune
disease, lupus, multiple sclerosis, transplant/graft rejection,
conditions (e.g. asthma, allergic rhinitis or atopic dermatitis). The
nucleic acids of the invention are also useful for down-regulating the
conditions (e.g. asthma, allergic rhinitis or atopic dermatitis).
Conditions (e.g. asthma, allergic Rhinitis) are aliagnostic tool to examine genetic
drifts and mutations within diseased cells or to detect the presence of a
target RNA in a cell. The present RNA sequence represents a human IKK-
gamma substrate sequence.
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allergy; asthma; allergic rhinitis; atopic dermatitis; Human IKK-gamma;
                                                                                                                                                                                                                                                                                                                       Novel enzymatic nucleic acid that down-regulates expression of neurite growth inhibitor receptor, prostaglandin D2 receptor, IkappaB kinase or protein kinase PKR genes, for treating cancer and inflammatory disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antisense oligonucleotide, neurite growth inhibitor; NOGO; prostaglandin D2 receptor; PTGDR; IkappaB kinase; IKK; protein kinase PKR; cerebrovascular accident; central nervous system injury; CNS injury; spinal cord injury; cancer; melanoma; lymphoma; glioma; inflammatory disease; rheumatoid arthritis; restenowis; asthma; Crohn's disease; diabbtes; obesity; autoimmune disease; lupus; multiple sclerosis; transplant rejection; graft rejection; ischaemia; reperfusion; glomerulonephritis; sepsis;
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                                                                                                                                                                                                                                                                                                                                                                                    Claim 59; SEQ ID NO 1837; 317pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human IKK-gamma substrate sequence #813.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            265
                                                                                                                                                                    05-APR-2001; 2001US-00827395.
29-MAY-2001; 2001US-0294412P.
28-AUG-2001; 2001US-0315315P.
                                                                                                                                       03-APR-2002; 2002WO-US010512
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                                                                                                                                                                                                                                 (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          249 TGGAGGCCGCGGTCGGC
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                                                                                                                                                                                                                                                                                             WPI; 2003-058513/05.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
                                                                           WO200281628-A2
                   substrate; ds.
                                                Unidentified.
                                                                                                        17-OCT-2002
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The invention comprises nucleic acids (e.g. antisense oligonucleotides)
that down regulate the expression or inhibit the function of a receptor
for a neurite growth inhibitor, NOGO, prostaglandin DZ receptor (PTGDR),
KappaB kinase (IKK), or protein Kinase PKR. The nucleic acids of the
invention are useful for treating: cerebrovascular accident, central
convention are useful for treating: cerebrovascular accident, central
convention are useful for treating: cerebrovascular accident, central
convention are useful for treating: creebrovascular accident, central
convention are useful for treating: creebrovascular accident, central
convention are useful for treating: creebrovascular accident, central
convention actions), inflammatory disease, diabetes, obesity, autoimmune
central reperfusion injury, glomerulonephritis, sepsis, and allergic
conditions (e.g. asthma, allergic rhinitis or atopic dermatitis). The
conditions (e.g. asthma, allergic rhinitis or atopic dermatitis). The
cutofic acides of the invention are also useful for down-regulating the
expression of a target gene and as a diagnostic tool to examine genetic
charges RNA in a cell. The present RNA sequence represents a human IKK-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
                                                                                                                                                                                     Novel enzymatic nucleic acid that down-regulates expression of neurite growth inhibitor receptor, prostaglandin D2 receptor, IkappaB kinase or protein kinase PKR genes, for treating cancer and inflammatory disease.
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NCH ribozyme; G-cleaver ribozyme; Zinzyme; DNAzyme; amberzyme; Inozyme;
halrpin ribozyme; tissue rejection; graft rejection; leukaemia.
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0
          Fosnaugh K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3.1%; Score 13.8; DB 1; Length 17; 76.5%; Pred. No. 3.4e+02; Ive 2; Mismatches 2; Indels
Haeberli P, Mcswiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 17 BP; 1 A; 6 C; 8 G; 0 T; 2 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                          Claim 59; SEQ ID NO 1836; 317pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human GRID mRNA substrate sequence #129.
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(BLAT/) BLATT L.
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                                                                                                                                                                                              leukemia
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                                    (JARV/)
                                                                                 (ELLI/)
                                               (CARL/)
(MCSW/)
                                                                       (HAMB)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 344
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       88;
HCV
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                                                                                                                                                                                                                                                                         The invention relates to a nucleic acid molecule that down-regulates expression of Grb2-related with insert domain (GRID) gene, e.g. a hammerhead ribozyme, NCH ribozyme, G-cleaver ribozyme, Zinzyme, DNAzyme, amberzyme, Inozyme or hairpin ribozyme. Also include are a mammalian cell call by contacting the cell with the novel nucleic acid molecule, reducing GRID activity in a cell by contacting the norel muth the novel nucleic acid molecule, treating a patient having a condition associated with the level of GRID cene by contacting the cell with the novel nucleic acid molecule, cleaving RNA of a GRID gene by contacting the cell with the novel nucleic acid molecule, newpression vector comprising a nucleic acid sequences (encoding at least the novel nucleic acid molecule in a manner that allows its expression), a mammalian cell including the expression vector and an enzymatic nucleic acid molecule that cleaves RNA derived from a GRID gene. The nucleic acid molecule is useful for treating a condition associated with the level of GRID, e.g. tissue/graft rejection and leukaemia. The present sequence is a target region for the enzymatic nucleic acide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                                                                                                          New nucleic acid molecule that down-regulates expression of Grb2-related with insert domain (GRID) gene, useful for treating a condition associated with the level of GRID, e.g. tissue/graft rejection and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; 88; GRID; Grb2-related with insert domain; hammerhead ribozyme;
NCH ribozyme; G-cleaver ribozyme; Zinzyme; DNAzyme; amberzyme; Inozyme;
hairpin ribozyme; tissue rejection; graft rejection; leukaemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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0
                                                                                                                                             Ellis JH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3.1%; Score 13.8; DB 1; Length 17; 88.2%; Pred. No. 3.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2; Indels
                                                                                                                                             Hamblin PA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 17 BP; 3 A; 5 C; 8 G; 0 T; 1 U; 0 Other,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human GRID mRNA substrate sequence #330.
                                                                                                                                             Mcswiggen J,
                                                                                                                                                                                                                                                      Claim 4; SEQ ID NO 129; 74pp; English.
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0
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                      23-FEB-2001; 2001US-00792818
                                               10-FEB-2000; 2000US-0181594P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         201 CTCCCGGGGACCTGCGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                             Jarvis T, Carlowitz IV,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity 88.2
les 15; Conservative
                                                                               CARLOWITZ I V.
MCSWIGGEN J.
HAMBLIN P A.
                                                                                                                                                                     WPI; 2003-829646/77
                                                                                             (MCSW/) MCSWIGGEN (HAMB/) HAMBLIN P (ELLI/) ELLIS J H.
                                                                       JARVIS T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US2003134806-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
17-JUL-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            03-JUN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17-JUL-2003
                                                                                                                                                                                                                                leukemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADM54055;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                     JARV/)
                                                                                 CARL/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
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The invention relates to a nucleic acid molecule that down-regulates
expression of Grb2-related with insert domain (GRID) gene, e.g. a
capression of Grb2-related with insert domain (GRID) gene, e.g. a
hammerhead ribozyme, NGH ribozyme, G-cleaver ribozyme, Zinzyme, Dinzyme, Dinzyme, Dinzyme, Dinzyme, Dinzyme, Dinclude are a mammalian cell
captuding the novel nucleic acid molecule, reducing GRID activity in a
cell by contacting the cell with the novel nucleic acid molecule,
creating a patient having a condition associated with the level of GRID
(e.g. tissue/graft rejection or leuksemia) by contacting the cell with
contacting the cell with the novel nucleic acid molecule, an expression
contacting the cell with the novel nucleic acid molecule, an expression
contacting the cell with the novel nucleic acid molecule, an expression
contacting the cell with the novel nucleic acid molecule, an expression
contacting the cell with the novel nucleic acid molecule in a manner that allows its expression, a
mammalian cell including the expression vector and an enzymatic nucleic
cacid molecule that cleaves RNA derived from a GRID gene. The nucleic acid
cacid molecule for treating a condition associated with the level of
GRID, e.g. tissue/graft rejection and leukaemia. The present sequence is
cacid molecule for the enzymatic nucleic acide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New nucleic acid molecule that down-regulates expression of Grb2-related with insert domain (GRID) gene, useful for treating a condition associated with the level of GRID, e.g. tissue/graft rejection and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gape
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    enzymatic nucleic acid, RNA cleavage; hepatitis C virus; HCV; infection; type I interferon; DNAzyme.
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                                                                                                                                                                                                                                                                                                                              Ellie JH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3.1%; Score 13.8; DB 1; Length 17; 88.2%; Pred. No. 3.46+02; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                              Hamblin PA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 17 BP; 3 A; 4 C; 9 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                              Mcswiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 4; SEQ ID NO 330; 74pp; English.
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10-FEB-2000; 2000US-0181594P.
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ID ADI85884 standard; RNA; 17
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                                                                                                                                                                                                                                                                                                                              Carlowitz IV,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity 88.2
Matches 15, Conservative
                                                                         JARVIS T.
CARLOWITZ I V.
MCSWIGGEN J.
HAMBLIN P A.
ELLIS J H.
                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2003-829646/77.
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US09436060A-16.rng.sl

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Claim 42; Page 83; 155pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   07-SEP-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAV35627;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           syndrome.
                                                                                                                                                                                                                                                                                                                                                                                        RESULT 346
AAV35627/c
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                                                                                                                                        Enzymatic nucleic acid molecules which specifically cleave RNA derived from hepatitis C virus (HCV), useful for the treatment of HCV infections, especially in combination with type I interferon therapy.
                                                                                                                                                                                                                                 The invention relates to an enzymatic nucleic acid molecule which specifically cleaves RNA derived from hepatitis C virus (HCV), in which the binding arms of the enzymatic nucleic acid molecule comprises sequences complementary to any of the defined substrate sequences given in the specification. The nucleic acid molecule may be administered for the treatment of HCV infections, especially in combination with type I interferons. The present sequence represents a HCV DNAzyme substrate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Maize, corn, Zea mays, delta-9 desaturase, GBSS, target, substrate, granule bound starch synthase; hammerhead ribozyme; hairpin ribozyme; modulation; gene expression; transgenic plant; cleavage; canola plant; caffeine synthesis; coffee plant; nicotine production; tobacco; fruit ripening; flower pigmentation; lignin production; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Skokut TA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ribozyme which modulates plant gene expression - preferably modulates expression of DELTA-9 desaturase or granule bound starch synthase in
                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Granule bound starch synthase hairpin substrate SEQ ID NO:571.
                                                                                                                                                                                                                                                                                                                                                                                                         3.1%; Score 13.8; DB 1; Length 17; 88.2%; Pred. No. 3.48+02; ive 0; Mismatches 2; Indels
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                                                                               Pavco PA, Macejack
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Merlo DJ;
                                                                                                                                                                                                                                                                                                                                                                           Sequence 17 BP; 3 A; 11 C; 2 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                       Claim 1; SEQ ID NO 3130; 198pp; English
                                                                               Roberts E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TGGGAGGGGTGGTGGCC 36
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                                                                                                                                                                                                                                                                                                                                                                                                                                         15; Conservative
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Folkerts O,
                                                                               Blatt L, Mcswiggen J,
   MCSWIGGEN J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1997-202224/18.
                ROBERTS E. PAVCO P A.
                                                                                                            WPI; 2004-031273/03
                                                MACEJACK D.
                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 maize or canola.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                12-JUL-1996;
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Young SA,
                                                                                                                                                                                                                                                                                                                                                sequence.
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(DOWC ) I
                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local
                                (PAVC/)
(MACE/)
   (MCSM/)
                   (ROBE/)
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The present invention describes an enzymatic nucleic acid molecule (I) with RNA cleaving activity, which modulates the expression of a plant gene. Also described is a gene comprising a cDNA sequence encoding maize Delta-9 desaturase. (I) can be used to modulate expression of a gene, preferably Delta-9 desaturase or a granule bound starch synthase (GBSS) gene, in a plant (preferably a maize or canola plant). (I) can be used to modulate caffeine synthesis in a coffee plant, nicotine production in a tobacco plant, fruit ripening processes in an apple, tomato, pear, plum or peach plant, flower pigmentation in a rose, petunia, chrysanthemum or marigold plant or lignin production in a tobacco, aspen, poplar or pine
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This exon specific primer used in the PCR amplification of a short stature associated sequence. The gene region corresponding to short stature has been identified as a region of approximately 500 kb in the PARI region of the X and Y chromosomes. The egenes in this region have been identified as candidates for the short stature gene. These genes were designated SHOX (also referred to as SHOX93 or HOX93), pET92 and SHOX (slox-like homeobox gene on chromosome three). The SHOX gene has two separate splicing sites resulting in two variations SHOXa and SHOXb. The specification provides sequences of SHOX (short stature homeobox-containing) genes SHOX ET92, SHOXb, SHOT and exons of the SHOX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homeobox domain; human growth gene; growth regulation; growth defect; turner's syndrome; short stature homeobox containing gene; short stature; SHOX; bone disease; osteoporosis; calcium regulation; PCR primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                Sequence 18 BP; 1 A; 9 C; 7 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                        1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 11; 84pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                     134 CGGCCTGCCGCCTTCCA 150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18
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97EP-00100583.
                                                                                                                                                                                                                                                                                                                         3.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CGGCCUGCCGCCGGCCA
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Best Local Similarity 82.4
Matches 14; Conservative
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genes as shown in AAV35610 to AAV35621 and protein sequences of the human growth protein transcription factor SHOXA, SHOXD and SHOT as shown AAW60573. The novel genes are responsible for human growth. Defects in the genes and cause short stature, e.g. Turner's syndrome. The products can be used to develop agents for the treatment of short stature or other human growth disorders. The products can also be used for providing a mitogenic effect on cells, e.g. for the treatment of bone diseases such as osteoporosis and diseases involved with disturbance in the bone calcium regulation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a novel probe for hybridising to a target nucleic acid and detecting gene polymorphisms. The probe comprises a base sequence that is complementary to the target nucleic acid and has at least one artificial mismatch and at least one original mismatch with respect to the target nucleic acid, where the artificial mismatch with original mismatch and effect in different base positions. The probe of the invention may be useful for detecting genetic variation by gene amplification, mutation-specific DNA sequencing or a mutation-specific DNA chip, preferably for detecting a polymorphism in exon 2 of HIA-DRB1. The current sequence is that of a probe of the invention which was used to detect a polymorphism in human HIA-DRB1 exon 2 DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   probe for detecting gene polymorphism, contains oligonucleotide is complementary to target sequence, has artificial mismatch with ct to target, and has original mismatch with respect to allelic
                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       polymorphism; genetic variation; exon 2; HLA-DRB1; human; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Probe 52 used to detect polymorphism in human HLA-DRB1 exon 2 DNA.
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0
                                                                                                                                                                                                                                3.1%; Score 13.8; DB 1; Length 18; 88.2%; Pred. No. 3.6e+02; ve 0; Mismatches 2; Indels
                                                                                                                                                                                              Sequence 18 BP; 0 A; 14 C; 4 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 18 BP; 2 A; 6 C; 9 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1; SEQ ID NO 52; 74pp; Japanese.
                                                                                                                                                                                                                                                                                                               332 CGGGGGCGAGGCGAGG 348
                                                                                                                                                                                                                                                                                                                                                                                                                                                   B
                                                                                                                                                                                                                                                    l Similarity 88.2%;
15; Conservative 0
                                                                                                                                                                                                                                                                                                                                   17 CGGGGCGGGGGGGGG 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (KOKU-) KOKUSAI SHIYAKU KK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
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                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           variant of target.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             sapiens
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Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADL88552;
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                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 347
                                                                                                                                                                                                                                                                        Matches
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This invention relates to a novel chemical sensor system method.

Specifically, it refers to an isolated nucleic acid molecule that encodes a receptor protein, which binds to chemicals that can stimulate the sense of taste or small for example. The present invention describes the manufacture of a chip that acts as a support to immobilise transfected calls expressing the receptor gene, such that this chip can be employed as a component of the chemical sensor model. Furthermore, this chip is useable as an artificial sensory organ where the chemical receptor contains an olfactory receptor the sensor can react to olfactory stimulation. Accordingly, these sensors are useful in the food industry contains environmental examination and disease diagnosis. Furthermore, such systems are automatable for high throughput applications under carrious conditions, even for differentiating optical isomers of R(-)-carvone easily. This oligonucleotide sequence is used to construct an odourant receptor DNA expression vector of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chemical sensor systems based on chemical receptors introduced into cells for immobilization onto support to form chip as component of sensor, useful in detecting stimuli e.g. taste and smell applicable in food
                                                                                                                                                                                                                                   DNA oligo to construct an odourant receptor expression vector SegID102.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gape
                                                                                                                                                                                                                                                                   ss; chemical sensor system; taste; smell; artificial sensory organ; olfactory stimulation; food industry; hygiene inspection; environmental examination; disease diagnosis; carvone.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sato T, Hirono J, Hamana H, Miyake M, Yoshikawa T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 18 BP; 5 A; 8 C; 4 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (NAAD-) NAT INST ADVANCED IND SCI & TECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 9; SEQ ID NO 102; 521pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     237 CCGAACCCCGCCTGGAG 253
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28-MAY-2002; 2002JP-00154239.
13-JUN-2002; 2002JP-00172412.
14-JAN-2003; 2003JP-00005175.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           28-MAY-2003; 2003WO-JP006719
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                88.2%;
262 CGGCCGGGGCTTCTCC
                        CGGCCCGGGCCTGCTCC
                                                                                                                           ADM42858 standard; DNA; 18
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                                                                                                                                                                                                 (first entry)
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                                                                                                                                                                                                                                                                                                                                               Synthetic.
                                   18
                                                                                                                                                               ADM42858;
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                                                                                         RESULT 348
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RESULT 349

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Gaps

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3.1%; Score 13.8; DB 1; Length 18; 88.2%; Pred. No. 3.6e+02; ive 0; Mismatches 2; Indels

Conservative

Query Match Best Local Similarity Matches 15; Conserva

AAZ07296/c

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The present invention relates to oligonucleotides that downregulate the expression of human Grb2-related with Insert Domain (GRID) gene. GRID is a T-cell co-stimulatory adaptor protein. The oligonucleotides are useful for modulating the expression of GRID, to treat conditions such as tissue/graft rejection and leukaemia. The oligonucleotides can also be administered in conjunction with other therapies such as radiation, chemotherapy and cyclosporin treatment. The present oligonucleotide was
                                                                                         Human, Grb2-related with Insert Domain; GRID; T-cell; co-stimulatory adaptor protein; tissue rejection; graft rejection; leukaemia; cytostatic; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human, Grb2-related with Insert Domain; GRID; T-cell; co-stimulatory adaptor protein; tissue rejection; graft rejection; leukaemia; cytostatic; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New nucleic acid(s) for regulating the Grb2-related with Insert D (GRID) gene comprises using antisense and enzymatic nucleic acid molecules such as hammerhead ribozymes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
3.0%; Score 13.4; DB 1; Length 17;
Best Local Similarity 93.3%; Pred. No. 3.6e+02;
Matches 14; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mcswiggen JA, Hamblin PA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human GRID NCH ribozyme substrate oligonucleotide #250.
                                                  Human GRID Amberzyme substrate oligonucleotide #124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 17 BP; 3 A; 3 C; 10 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 4; Page 87; 108pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  used to illustrate the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BB
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                                                                                                                                                                                                                                                                                                                                                                                                                                  24-FEB-2000; 2000US-0184594P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                23-FEB-2001; 2001WO-US005957.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Jarvis T, Von Carlowitz I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (RIBO-) RIBOZYME PHARM INC. (GLAX ) GLAXO GROUP LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15 CTCGCCTGCCGCT 1
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  27-JUN-2003 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-550088/61.
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                                                                                                                                                                                                                  Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The INVELLOR INTERIOR PROGREMS FLOW MINERS AND HIGHER TO COMPARE THE TRY COMPONENT GENERAL MINERS AND THE TRY OF MANY HUMAN CANCER. MUTATION OF UNDERGULATION OF THE VERONESTION TO AND THE VERONESTION TO CANCER. SUBSTANCES, I CANCER SHOPE EXPENSION IN CANCER. SUBSTANCES, I CANCER SHOPE TRY GENERAL MINERS AND THE MINERS AND THE TRY OF THE TRY OFTEN THE TRY OF THE TRY OFTEN THE TRY OF THE TRY OFTEN THE TRY OF THE TRY OF THE TRY OFTEN T
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to promoter regions from mouse and human telomerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mouse and human telomerase RNA gene promoters, useful for tumor specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                         Telomerase RNA; TR; promoter; cytotoxin; cancer; neoplasia; hTR; gene therapy; thymidine kinase gene; anticancer therapy; human; mutagenesis; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ;
0
                                                                                                                                                                                         Human telomerase RNA gene (hTR) promoter specific primer hll.
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                            BP.
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                         AAZ07296 standard; DNA; 38
                                                                                                                                    (first entry)
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                    22-OCT-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                  WO9938964-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29-JAN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    gene therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       05-AUG-1999.
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                                                                                                                                                                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Keith WN;
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ABL47224/c ID ABL472 XX AC ABL472 XX

RESULT 350

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Gaps

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Domain

Ellis JH;

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conditions selected from cancer, lymphoma, Ewing's sarcoma, melanoma, conditions selected from cancer, lymphoma, Ewing's sarcoma, melanoma, cumour angiogenesis, diabetic retinopathy, macular degeneration, necessitar angiogeneration, arthritis, psoriasis, vertuca vulgaris, angiofibroma of tuberous sclerosis, port-wine stains, Sturge where yindrome, Rippel-Trenaunay-Weber syndrome, Osler-Weber-rendu syndrome, Aippel-Trenaunay-Weber syndrome, Osler-Weber-rendu syndrome, apatient having a condition associated with the level of ERG, treating a patient having a condition associated with the level of ERG, by contacting cells of the patient with (I) under conditions suitable for the treatment. Leukaemia or tumour conditions suitable for the treatment. Leukaemia or tumour conditions suitable for the treatment. Leukaemia or tumour conjunction with one or more of other therapies such as radiation or conjunction with one or more of other therapies such as radiation or chemotherapy treatment. (I) is useful for reducing ENG activity in a coll, by contacting (I) the useful for dispance of advance call, by contacting (I) the useful for dispance of advance call of the expression of ERG, and as diagnostic tool to cation such as Mg2+. (I) is useful for specifically cateding genes that share homology with ERG gene or ERG fusion genes. Consumers of the stage of conditions and call of the specifically cargeting genes that share homology with ERG gene or ERG fusion genes.
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CYS4502A6; CYP2A6, chromosome 19; eteroid metabolism;
drug detoxification; xenobiotic detoxification; procarcinogen activation;
inflammation; asthma; habitual smoking.
 useful for treating cancer, diabetic retinopathy, macular degeneration, arthritis, psoriasis, verruca vulgaris and Sturge Weber syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                enzymatic nucleic acid molecules which regulate expression of ERG, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human cytochrome P4502A6 (CYP4502A6) promoter - polymorphic region 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /*tag= a
/standard_name= "Single nucleotide polymorphism"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3.0%; Score 13.4; DB 1; Length 17; 93.3%; Pred. No. 3.6e+02; tive 0; Mismatches 1; Indels
                                                                                             relates to a nucleic acid molecule (I) which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 17 BP; 1 A; 5 C; 6 G; 0 T; 5 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   related PCR primers of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers replace(10, C)
                                                        Claim 4; Page 93; 149pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ВР
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    124 GGAAAAGCCTCGGCC 138
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAL44028 standard; DNA; 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200194633-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 353
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                                                                                                                                                                                                                                                                               The present invention relates to oligonucleotides that downregulate the expression of human Gry2-related with Insert Domain (GRID) gene. GRID is a T-cell co-stimulatory adaptor protein. The oligonucleotides are useful for modulating the expression of GRID, to treat conditions such as tissue/graft rejection and leukaemia. The oligonucleotides can also be administered in conjunction with other therapies such as radiation, chemotherapy and cyclosporin treatment. The present oligonucleotide was
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel polynucleotide which down regulates expression of Ets-related gene,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             neovascular glaucoma, myopic degeneration, arthritis, verruca vulgaris, angiofibroma of tuberous sclerosis, port-wine stain, wound healing; Sturge Weber syndrome, Kippel-Trenaunay-Weber syndrome, leukaemia, 88;
                                                                                                                                                                New nucleic acid(s) for regulating the Grb2-related with Insert Domain (GRID) gene comprises using antisense and enzymatic nucleic acid molecules such as hammerhead ribozymes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ophthalmological, antiarthritic, antipsoriatic, virucide, osteopathic, vulnerary, cancer, lymphoma; Ewing's sarcoma; melanoma, psoriasis, tumour angiogenesis, diabetic retinopathy, macular degeneration;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Osler-Weber-rendu syndrome, leukaemia; osteoporosis; DNAzyme; inozyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Randi AM
                                                                                           Ellis JH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; hammerhead ribozyme; cytostatic; antitumour; antidiabetic;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3.0%; Score 13.4; DB 1; Length 17; 93.3%; Pred. No. 3.6e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1; Indels
                                                                                           Mcswiggen JA, Hamblin PA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 17 BP; 3 A; 4 C; 9 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human ERG DNAzyme target sequence Seq ID No 1481.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
                                                                                                                                                                                                                                            Claim 4; Page 67; 108pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                used to illustrate the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABK18834 standard; RNA; 17 BP
24-FEB-2000; 2000US-0184594P.
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                                                                                           Von Carlowitz I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RIBOZYME PHARM INC
GLAXO GROUP LTD.
                                     (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                14; Conservative
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                                                      (GLAX ) GLAXO GROUP LTD
                                                                                                                               WPI; 2001-550088/61.
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                                                                                           Jarvis T,
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                                                                                                                                The invention comprises the identification of genetic polymorphisms in the human cytochrome P4502A6 (CYP4502A6 or CYP2A6) gene. The human cytochrome P4502A6 (CYP4502A6 or CYP2A6) gene. The human cytochrome P4502A6 gene is located on chromosome 19 and encodes an enzyme that plays a role in the metabolism of steroids, the detoxification of drugs and xenobiotics, and the activation of procarcinogens. The P4502A6 polymorphisms identified in the invention are useful for evaluating an individual's risk of developing asthma or an individual's propensity for cigarette consumption. The P4502A6 DNA sequences of the invention are useful for identifying individuals having a polymorphic genotype and to screen individuals for altered metabolism for cytochrome P4502A6 substrates. The P4502A6 DNA sequences of the invention are also useful for identifying individuals who are at risk from inflammation, asthma, habitual smoking and diseases that result from environmental or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; ds; single nucleotide polymorphism; SNP; cytochrome; P4502A6; CYP4502A6; CYP2A6; chromosome 19; steroid metabolism; drug detoxification; xenobiotic detoxification; procarcinogen activation;
                                                                                                                                                                                                                                                                                            occupational exposures to dangerous substances. The present DNA sequence represents a polymorphic region of the promoter for the human cytochrome P4502A6 gene
                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human cytochrome P4502A6 (CYP4502A6) promoter - polymorphic region 4.
                                                            isolated polynucleotide, useful to screen individuals for asthma,
                                                                         inflammation and susceptibility to habitual smoking, comprises base variation from that of known human cytochrome P4502A6 sequence.
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                                                                                                                                                                                                                                                                                                                                                Sequence 17 BP; 4 A; 5 C; 1 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  inflammation; asthma; habitual smoking.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                              Claim 35; Page 27; 48pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               replace(10, T)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAL44029 standard; DNA; 17 BP
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                                                                                                                                                                                                                                                                                                                                                                                                  14; Conservative
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                                     WPI; 2002-566448/60
                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
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            Guida M, Hall J;
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variation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAL44029;
                                                                                                                                                                                                                                                                                                                                                                           Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            represents a polymorphic region of the promoter for the human cytochrome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                     New isolated polynucleotide, useful to screen individuals for asthma, inflammation and susceptibility to habitual smoking, comprises base variation from that of known human cytochrome P4502A6 sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 17 BP; 4 A; 6 C; 1 G; 6 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                       Claim 1; Page 27; 48pp; English
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94US-00245466.
94US-00291932.
96US-00777916.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 93.33
Matches 14; Conservative
WPI; 2002-566448/60.
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15-AUG-1994;
23-DEC-1996;
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Homo sapiens
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                                                                  Mcswiggen J;
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                                                                                                                                                                                The invention describes an enzymatic nucleic acid molecule (I) which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B (NFRS), where (I) is an inozyme, zinzyme, G-cleaver or amberzyme configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulating REL-A activity in a cell, for treating a patient having a condition associated with the level of REL-A. (I) is useful for cleaving RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially MG-2+. The enzymatic and antisense nucleic acid molecules are useful for treating breast, lung, prostate, colorectal, brain, oesophagel, stoomach, bladder, pancreatic, cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or multidrug resistant cancer. The method involves use of other drug therapies such as monoclonal antibodies, REL-A-specific inhibitors or chemotherapy including paclitaxel, docetaxel, cisplatin, methotrexate, cyclophosphamide, doxorubin, fluorouzacil carboplatin, methotrexate, gemiclabine or radiation therapy. The enzymatic and antisense nucleic acid molecules are also useful for treating inflammatory disease such as rheumatoid arthritis, restenosis, asthma. Crohn's disease, diabetes, obesity, autoimmune disease, lupus, multiple sclerosis, transplant/graft rejection, gene therapy applications, asthma. Control inflammatory bowel disease or infection. The sequence represents the substrate of a novel enzymatic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                      Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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enzymatic nucleic acid, H-Ras, N-Ras, HIV, cytostatic, anti-HIV,
anti-rheumatic, cancer, AIDS, 88.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3.0%; Score 13.4; DB 1; Length 17; 73.3%; Pred. No. 3.6e+02; tive 3; Mismatches 1; Indel8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 17 BP; 0 A; 9 C; 4 G; 0 T; 4 U; 0 Other;
                                                    Stinchcomb DT, Mcswiggen J, Draper KG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human HIV DNA zyme substrate #13.
                                                                                                                                                          Claim 3; Page 38; 72pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        132 CTCGGCCTGCCGCT 146
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STINCHCOMB D T. MCSWIGGEN J.
                                                                             WPI; 2003-340953/32.
                          DRAPER K G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity
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(STIN/)
                          (DRAP/)
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ABZ66556
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The invention relates to a novel short interfering RNA (siRNA) nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HER2, K-Ras, H-Ras, N-Ras, human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule are useful, and anti-rheumatic activity. The nucleic acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung, prostate, bladder, or pancreatic cancer, and HIV infection, and AIDS. The sequences shown in ABZ59889 - ABZ62216, ABZ64544 - ABZ65531, ABZ6520 - ABZ65224, ABZ65530 - ABZ65216, Human
                                                                                                                                                                                                                                                                                                                                                                  Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 3.0%; Score 13.4; DB 1; Length 17; Best Local Similarity 80.0%; Pred. No. 3.6e+02; Matches 12; Conservative 2; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 123; Page 157; 185pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human HIV amberzyme substrate #19.
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06-JUN-2001; 2001US-0296249P.
10-SEP-2001; 2001US-0318471P.
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06-JUN-2001; 2001US-0296249P.
10-SEP-2001; 2001US-0318471P.
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                                                                                                                    (RIBO-) RIBOZYME PHARM INC.
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(RIBO-) RIBOZYME PHARM INC.
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                                                                                                  infection.
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                                                      Novel
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                                                                  The invention relates to a novel short interfering RNA (siRNA) nucleic acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HER2, K-Ras, H-Ras, N-Ras, human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-theumatic activity. The nucleic acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung, prostate, bladder, or pancreatic cancer, and HIV infection, and AIDS. The sequences shown in ABZ56859 - ABZ665216, ABZ66531, ABZ66520 - ABZ66524, ABZ66530 - ABZ66554 or the human ribozymes of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nucleic acid molecule; Hepatitis C virus; HCV; Hepatitis B virus; HBV; RNA stability; RNA expression; RNA synthesis; antisense; enzymatic nucleic acid, harmerihead ribozyme; DNAzyme; inozyme; amberzyme; G-cleaver ribozyme; decoy molecule; aptamer; HBV reverse transcriptase; Enhancer I region; viral replication; degenerative, disease state, HBV infection; HCV infection; cirrhosis; liver failure; hepatocellular carcinoma; hepatotropic; cytostatic; virucide; antiinflammatory; substrate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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  HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences.
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                                                                                                                                                                                                                                                                                                                                                                               3.0%; Score 13.4; DB 1; Length 17;
80.0%; Pred. No. 3.6e+02;
tive 2; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            HCV minus strand DNAzyme substrate sequence #1361.
                                                                                                                                                                                                                                                                                                                                          Seguence 17 BP; 3 A; 6 C; 6 G; 0 T; 2 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Morrissey D,
                                         Claim 123; Page 158; 185pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mcswiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2001US-00877478.
2001US-0296876P.
2001US-0335059P.
2001US-0337055P.
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BLATT L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                         12; Conservative
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Roberts E;
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MACEJAK D.
MCSWIGGEN J.
MORRISSEY D.
                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity
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LEE P.
DRAPER K.
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08-JUN-2001; 2
08-JUN-2001; 2
24-OCT-2001; 2
05-DEC-2001; 2
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(BLAT/)
(MACE/)
(MCSW/)
(MORR/)
(PAVC/)
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(ROBE/)
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The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HCV) or the synthesis, expression and/or stability of Hepatitis C virus (HCV) or and enzymatic nucleic acids such as hammerhead ribozymes. DNAzymes, and enzymes, zinzymes, amberzymes, and G-cleaver ribozymes. DNAzymes, or are nucleic acid decoy molecules and aptamers that bind to HBV reverse transcriptase primer sequences, as well constructed and/or HBV reverse transcriptase primer sequences, as well as oligonucleotides that specifically bind the Enhancer I region of HBV CC DNA. The nucleic acids may be used to modulate the expression of HBV compounds and/or potential therapies directed against HBV, and compounds compounds and/or potential therapies directed against HBV, and compounds confidence acids the average of the compounds and disease states related to HBV and HCV infection, replication and gene compounds and disease states related to HBV and HCV infection, replication and gene compounds and sequence represents a substrate for one of the HCV DNAZYME or minus strand DNAZYME sequences disclosed in the present
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                                                                    Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus
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                                                                                                                                                                                                                                 Claim 1; Page 299; 387pp; English.
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08-JUN-2001; 2001US-0296876P.
24-OCT-2001; 2001US-0335059P.
05-DEC-2001; 2001US-0337055P.
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WPI; 2003-229207/22.
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Best Local Similarity
Matches 14; Conserv
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17-SEP-2001; 2001FR-00011980.
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                                                                                                                                                                                                                 The present invention relates to nucleic acid molecules which modulate
the synthesis, expression and/or stability of Hepatitis C virus (HCV) or
Hepatitis B virus (HBV) RNA. The nucleic acid molecules include antisense
and enzymatic nucleic acids such as hammerhead ribozymes, DNAzymes,
inozymes, zhrzymes, amberzymes, and G-cleaver ribozymes. Also disclosed
are nucleic acid decoy molecules and aptamers that bind to HBV reverse
transcriptase and/or HBV reverse transcriptase primer sequences, as well
as oligonucleotides that specifically bind the Enhancer I region of HBV
DNA. The nucleic acids may be used to modulate the expression of HBV
compounds and/or potential therapies directed against HBV, and compounds
that modulate the expression and/or replication of HCV. The compounds and
methods of the invention are useful for the treatment of degenerative and
carcinoma. The present sequence represents a substrate for one of the HCV
conversion.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              tumour suppression; tumour reversion; apoptosis; virus resistance; cytostatic; virucide; neuroprotective; nootropic; neuroleptic; probe; primer; PCR; gene chip; antisense; viral disease; tumour; cell degeneration; cancer; Alzheimer's disease; schizophrenia; ds; human.
                                                                                                                                                    Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human tumour suppression/reversion-related DNA sequence SeqID1134.
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80.0%; Pred. No. 3.6e+02;
tive 2; Mismatches 1; Indels
                                                                                                Pavco P,
                                                                                                Mcswiggen J, Morrissey D,
                                                                                                                                                                                                                                                                                                                                                                                                                                Seguence 17 BP; 1 A; 7 C; 7 G; 0 T; 2 U; 0 Other;
                                                                                                                                                                                                Claim 1; Page 250; 387pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2 ACCUGCGGCGGCUCG 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15-APR-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches 12; Conservative
                                                                                               Macejak D,
Roberts E;
BLATT L.
MACEJAK D.
MCSWIGGEN J.
MORRISSEY D.
                                                                                                                                WPI; 2003-229207/22
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
                                          PAVCO P.
LEE P.
DRAPER K.
                                                                           ROBERTS E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO2003025177-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               27-MAR-2003
                                                                                                                                                                           infection.
                                                                                                           Draper K,
                                                                                               Blatt L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADI48631;
                    (MCSW/) N
(MORR/) N
(PAVC/) I
(LEEP/) I
(DRAP/) I
(ROBE/) H
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
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This invention relates to novel isolated nucleic acid sequences involved in the phenomena of tumour suppression, tumour reversion, apoptosis and/or resistance to viruses. The invention may be useful for the anotropic or neuroleptic activity. The DNA sequences may be useful as probes and primers for detecting, indentifying, quantifying and/or probes and primers for detecting, indentifying, quantifying and/or complitying nucleic acid, for example as one component of a gene chip, in vitro as antisense reagents and for production of recombinant polypeptides. The invention may therefore be useful for preparation of polypeptides. The invention and/or treatment of viral diseases that are characterised by development of tumours or cell degeneration, specifically cancer but also Alzheimer's disease and schizophrenia. The present sequence is that of a nucleic acid sequence of the invention.

Once: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                            New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human, 89, GRID, Grb2-related with insert domain, hammerhead ribozyme,
NCH ribozyme, G-cleaver ribozyme, Zinzyme, DNAzyme, amberzyme, Inozyme,
hairpin ribozyme, tissue rejection, graft rejection, leukaemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 3.0%; Score 13.4; DB 1; Length 17; Best Local Similarity 93.3%; Pred. No. 3.6e+02; Matches 14; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 17 BP; 8 A; 3 C; 5 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            at ftp.wipo.int/pub/publishedpct_sequences
                                                                                                                                                                                                                                                                                                                                                                   Disclosure; SEQ ID NO 1134; 30pp; French
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human GRID mRNA substrate sequence #855.
                                                                       Ξ
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(MOLE-) MOLECULAR ENGINES LAB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10-FEB-2000; 2000US-0181594P.
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MCSWIGGEN J.
HAMBLIN P A.
ELLIS J H.
                                                                                                                                                                                                                                                                                   and transfected cells.
                                                                       relerman A, Amson R,
                                                                                                                                           WPI; 2003-313354/30.
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(MCSW/)
(HAMB/)
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Page 184

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The invention relates to a nucleic acid molecule that down-regulates carpession of Grb2-related with insert domain (GRID) gene, e.g. a hammerhead ribozyme, NCH ribozyme, Alizyme, Zhizyme, DNAzyme, amberzyme, Inozyme or hairpin ribozyme. Also include are a mammalian cell calluding the novel nucleic acid molecule, reducing GRID activity in a cell by contacting the cell with the novel nucleic acid molecule, reducing GRID activity in a cell by contacting the cell with the level of GRID (e.g. tissue/graft rejection or leukaemia) by contacting the cell with the novel nucleic acid molecule, cleaving RNA of a GRID gene by contacting the cell with the novel nucleic acid molecule, an expression vector comprising a nucleic acid sequences (encoding at least the novel nucleic acid molecule in a manner that allows its expression, a mammalian cell including the expression vector and an enzymetic nucleic acid molecule is useful for treating a condition associated with the level of GRID, e.g. tissue/graft rejection and leukaemia. The present sequence is a target region for the enzymatic nucleic acide that region for the enzymatic nucleic acide target region for the enzymatic nucleic acide acides of the invention.
                                                New nucleic acid molecule that down-regulates expression of Grb2-related with insert domain (GRID) gene, useful for treating a condition associated with the level of GRID, e.g. tissue/graft rejection and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      enzymatic nucleic acid; RNA cleavage; hepatitis C virus; HCV; infection; type I interferon; DNAzyme.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3.0%; Score 13.4; DB 1; Length 17; 93.3%; Pred. No. 3.6e+02; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 17 BP; 3 A; 3 C; 10 G; 0 T; 1 U; 0 Other;
                                                                                                                                 Claim 4; SEQ ID NO 857; 74pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HCV DNAzyme substrate sequence #3638.
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Best Local Similarity 93.3%,
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MCSWIGGEN J.
ROBERTS E.
PAVCO P A.
MACEJACK D.
                 WPI; 2003-829646/77
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                                                                                                   leukemia.
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(ROBE/)
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(MACE/)
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ID ADI8
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Pavco PA,

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from hepatitis C virus (HCV), useful for the treatment of HCV infections, especially in combination with type I interferon therapy.
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                                                                        The invention relates to an enzymatic nucleic acid molecule which specifically cleaves RNA derived from hepatitis C virus (HCV), in which the binding arms of the enzymatic nucleic acid molecule comprises sequences complementary to any of the defined substrate sequences given in the specification. The nucleic acid molecule may be administered for the treatment of HCV infections, especially in combination with type I interferons. The present sequence represents a HCV DNAzyme substrate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         / cdbase= OTHER
/mode= "Sugar-phosphate backbone has been replaced by
peptide backbone"
                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Peptide nucleic acid; PNA; cancer; telomerase; probe; hybridisation;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Peptide nucleic acid 4, targeted to mammalian telomerase.
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                                                                                                                                                                                                           Sequence 17 BP; 2 A; 7 C; 6 G; 0 T; 2 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Corey D,
                                            Claim 1; SEQ ID NO 3638; 198pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Piatyszek MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cocation/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 9; Page 59; 76pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                            ВЪ.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       diseases, also for diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       97WO-US005931
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(first entry)
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Best Local Similarity
Matches 14; Conserv
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modified_base
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Gaps

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Indels

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Mismatches

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Conservative
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12-MAY-1998
  13;
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                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic
                                                                           13
                                                                                                                                                                                                                AAT89225;
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AAZ08815/c
  Matches
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mammalian telomerase and as inhibitors of telomerase activity, especially
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New peptide nucleic acids hybridising to mammalian telomerase RNA - used to inhibit telomerase, for treating tumours and other proliferative diseases, also for diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /*tag= a
/mod base= OTHER
/note= "Sugar-phosphate backbone has been replaced by a
peptide backbone"
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                                                           Revised record issued on 21-OCT-2004 : Correction to feature table key
                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    nucleic acid; PNA; cancer; telomerase; probe; hybridisation;
                                                                                                                                                                             ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Peptide nucleic acid 11, targeted to mammalian telomerase.
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                                                                                                                                      Length 13;
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                                                                                                                                                                           0; Indels
                                                                                                                                  Score 13; DB 1; Le
Pred. No. 2.8e+02;
0; Mismatches 0;
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                                                                                               Sequence 13 BP; 5 A; 1 C; 4 G; 3 T; 0 U; 0 Other
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                                                                                                                    2.9%; Scor.
100.0%; Pre
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                      in the treatment of cancer
                                                                                                                                                                                                                                                                                                                                                       AAT89236 standard; DNA; 13
                                                                                                                                                                                                                                                                                                                                                                                                                                      (revised)
(first entry)
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                                                                                                                                                                                                                    42 TIGICIAACCCIA 54
                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                         13 rrercraaccera 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Shay JW, Wright WE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1997-512647/47.
                                                                                                                                  Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           inhibitor; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        09-APR-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO9738013-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                    21-OCT-2004
12-MAY-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic
                                                                                                                                                                                                                                                                                                                                                                                               AAT89236;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Peptide
                                                                                                                                                                                                                                                                                                                  RESULT 364
                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                      AAT89236/C
AAC
AAT89
XX
AAC
AAT89
XX
AAT89236/C
DT 21-OC
DT 12-M9
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Best Local Similarity

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New peptide nucleic acids hybridising to mammalian telomerase RNA - used to inhibit telomerase, for treating tumours and other proliferative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note= "Sugar-phosphate backbone has been replaced by a peptide backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Revised record issued on 21-OCT-2004 : Correction to feature table key
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gape
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Peptide nucleic acid; PNA; cancer; telomerase; probe; hybridisation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                                                                                                                                                   Peptide nucleic acid 1, targeted to mammalian telomerase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2.9%; Score 13; DB 1; Length 13; 00.0%; Pred. No. 2.8e+02; ve 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Norton JC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 13 BP; 3 A; 1 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Corey D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Piatyszek MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /*tag= a
/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 9; Page 59; 76pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      diseases, also for diagnosis.
                                                                                                                                                                                                   AAT89225 standard; DNA; 13
                                                                                                                                                                                                                                                                                                                (revised)
(first entry)
44 GICTAACCCTAAC 56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               46 CTAACCCTAACTG 58
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Matches 13; Conservative
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                                                     GTCTAACCCTAAC 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1997-512647/47.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Key
modified_base
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The present sequence represents a peptide nucleic acid molecule which hybridises to the mRNA component of mammalian telomerase, and inhibits telomerase activity. Telomerase is a ribonucleoprotein enzyme that synthesizes one strand of the telomeric DNA, using as a template an ill nucleotide sequence contained within the RNA component of the enzyme. The invention relates to PNA molecules having a sequence of no more than 25 concers, which include the sequence of TTAGG. The uncharged nature of the PNA backbone increases the melting temperature of associating strands, increases the rate of association with targeted nucleic acids, and affords greater resistance of degradation by proteases or nucleases. The cancers, neoplasia, hyperplasia, neurodegenerative diseases. The deformance of degradation by proteases or nucleases. The cancers, neoplasia, hyperplasia, neurodegenerative diseases. The deformance of degradation with summodeficiency virus (HVV) infection/ANDS (acquired immunodeficiency virus (HVV) infection/ANDS (acquired immunodeficiency virus (HVV) infection/ANDS (acquired immunodeficiency syndrome) and associated pathologies, fungal infections, and other cyclocation of the activity, in combination with antineoplastic and other cyclocation of cyclotatic agents, antifungal agents, and other nucleotides. PNAs may be sectivity, in combination with antineoplastic and other cyclocation probes to detect or quantitate polynucleotides having a human telomerase and so triament length polymorphism (RFLP) pattern. PNAs are also useful as probes to detect the RNA component of a mammalian telomerase and as inhibitors of telomerase activity. The method of the present invention can be used to detected as cancerous conditions to be detected with increased confidence and pathological characteristics. The diagnostic and prognostic and proplastic cell or expresses telomerase activity and its RNA component immortal or mechads of the present invention can be used to detect an immortal or mechads of the present activity and its RNA 
                                                                                                                                                                                                                                                                                                 New peptide nucleic acid (PNA) compounds that inhibit telomerase activity in mammalian cells is useful as probes to detect the RNA component of a mammalian telomerase.
via glycine amino N through a methylenecarbonyl linker"
                                                                                                                                                                                                                            Corey DR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2.9%; Score 13; DB 1; Length 13;
100.0%; Pred. No. 2.8e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 13 BP; 3 A; 1 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                            Wright WE, Piatyszek MA, Shay JW, Norton JC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; Prec. ... 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                Claim 6; Col 71; 45pp; English.
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                                                                                                              97US-00838545
                                                                                                                                                   96US-00630019
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                                                                                                                                                                                       (TEXA ) UNIV TEXAS SYSTEM.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          46 CTAACCCTAACTG 58
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Matches 13; Conservative
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                                                                                                                                                                                                                                                                  WPI; 2000-292432/25.
                                                                                                            09-APR-1997;
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                                       US6046307-A
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                                                                          04-APR-2000
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AAA37594/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention describes an antitumour augmenting agent comprising a hybridisable oligonuclectide, as an effective component, contained at the template region of RNA site contained in telomerase. Also described
                                                                                                                                                 Human; RERF-LC-A1; hybridisation; antitumour augmenting drug; telomerase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Peptide nucleic acid; PNA; telomerase; ribonucleoprotein enzyme; cancer; inhibitor; neoplasia; neurodegenerative disease; aging; hyperplasia; AIDS; HIV; fungal infection; forensic identification; detect; tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           is an antitumour composition comprising the antitumour augmenting agent and an antitumour agent as effective components. The present sequence represents a hybridisable oligonucleotide for human RERF-LC-A1, used in the exemplification of the present invention. When the antitumour composition is administrated into a mammal, especially human, a sufficient antitumour effect is exhibited to cancer cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /*tag= a //rag= nucleic acid molecule, where N-(2-note= "Peptide nucleic acid molecule nucleotide bases
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New composition - for anti-tumor augmenting agent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Seguence 13 BP; 5 A; 1 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                            Human RERF-LC-Al hybridisable oligonucleotide.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 1; Page 6; 8pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                    (KURE ) KUREHA CHEM IND CO LTD
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 AAZ08815 standard; DNA; 13
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                                                                          (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1999-522625/44
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                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Key
misc_feature
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                                                                                                                                                                                                                                                                                                                                             10-FEB-1998;
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                                                                          01-NOV-1999
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                                                                                                                                                                                                                                                                                                       24-AUG-1999
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                                                                                                                                                                         cancer; ss.
                                                                                                                                                                                                             Synthetic.
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                                       AAZ08815
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XX AC AAA33
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The present sequence represents a peptide nuclear acts molecule which the present sequence represents a peptide nuclear acts molecule which telomerase activity. Telomerase is a ribonucleoprotein enzyme that telomerase activity. Telomerase is a ribonucleoprotein enzyme that telomerase activity. Telomerase is a ribonucleoprotein enzyme that concledide sequence contained within the RNA component of the enzyme. The nucleotide sequence contained within the RNA component of the enzyme. The invention relates to PNA molecules having a sequence of no more than 25 bases, which include the sequence GTTAGG. The uncharged nature of the PNA backbone increases the melting temperature of associating strands, increases the rate of association with targeted nucleic acids, and affords greater resistance of degradation by proteases or nucleases. The therapeutic PNAs may be used for treating diseases or nucleases. The therapeutic PNAs may be used for treating diseases conditions such as cancers, neoplasia, hyperplasia, neurodegenerative diseases, aging, human immunodeficiency virus (HIV) infection/AIDS (acquired immunodeficiency syndrome) and associated pathologies, fungal infections, and other cycloxic or cycostated by abnormal telomera metabolism or telomerase activity, in combination with antineoplastic and other cycloxic or cycostated by abnormal telomera metabolism or telomerase activity and other nucleotides having a human telomerase conditions to defect the RNA component of a mammalian telomerase and as probes to detect the RNA component of a mammalian telomerase and as probes to detect the RNA component of a mammalian telomerase and as inhibitors of telomerase activity. The method of the present invention allows cancerous conditions to be detected with increased confidence and or nathological characteries activity. The method of the present invention has an underlast and as an explanation of the present invention of hased on nathological characteries activity. The method of the present invention of hases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            , peptide nucleic acid (PNA) compounds that inhibit telomerase activity mammalian cells is useful as probes to detect the RNA component of a
Peptide nucleic acid, PNA; telomerase, ribonucleoprotein enzyme; cancer; inhibitor; neoplasia; neurodegenerative disease; aging; hyperplasia; AIDS; HIV; fungal infection; forensic identification; detect; tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note= "Peptide nucleic acid molecule, where N-(2-aminoethyl)glycine units are linked to nucleotide bases via glycine amino N through a methylenecarbonyl linker"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            based on pathological characteristics. The diagnostic and prognostic methods of the present invention can be used to detect an immortal or neoplastic cell or tumour tissue or cancer of any origin, provided the cell expresses telomerase activity and its RNA component
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      present sequence represents a peptide nucleic acid molecule which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Corey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Seguence 13 BP; 7 A; 1 C; 3 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Wright WE, Piatyszek MA, Shay JW, Norton JC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           a
"Peptide nucleic
                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
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                                                                                                                                                                  paternity testing; ss.
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New peptide nucleic acid (PNA) compounds that inhibit telomerase activity
                                                                                                                      Peptide nucleic acid; PNA; telomerase; ribonucleoprotein enzyme; cancer; inhibitor; neoplasia; neurodegenerative disease; aging; hyperplasia; AIDS; HIV; fungal infection; forensic identification; detect; tumour; paternity testing; ss.
                                                                                                                                                                                                                      /note= "Peptide nucleic acid molecule, where N-(2-aminoethyl)glycine units are linked to nucleotide bases via glycine amino N through a methylenecarbonyl linker"
                                                                                                                                                                                                                                                                                                                                                                                                   in mammalian cells is useful as probes to detect the RNA component of mammalian telomerase.
                                                                                                                                                                                                                                                                                                                                                          Piatyszek MA, Shay JW, Norton JC, Corey DR;
                                                                                                      PNA sequence #56 used to inhibit telomerase activity.
                                                                                                                                                                                           Location/Qualifiers
                                               AAA37598 standard; DNA; 13 BP.
                                                                                                                                                                                                                                                                                                   97US-00838545.
                                                                                                                                                                                                                                                                                                                     96US-00630019
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                                                                 AAA37598;
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                             RESULT 369
AAA37598/c
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The present sequence represents a peptide nucleic acid molecule which hybridises to the mRNA component of mammalian telomerase, and inhibits ctelomerase activity. Telomerase is a ribonucleoprotein enzyme that extended the telomeric DNA, using as a template an 11 structucities sequence contained within the RNA component of the enzyme. The sucleotide sequence contained within the RNA component of the enzyme. The nucleotide sequence contained within the RNA component of the enzyme. The concerns which include the sequence GTTAGG The uncharged nature of the PNA cackbone increases the melting temperature of associating strands, increases the rate of association with targeted nuclein acids, and affords greater resistance of degradation by proteases or nucleases. The charapeutic PNAs may be used for treating disease conditions such as cancers, neoplasia, hyperplasia, neurodegenerative diseases. The characterized by abnormal telomere metabolism or telomerase communication with antineoplastic and other cytcoxic or sectivity, in combination with antineoplastic and other cytcoxic or cytcostatic agents, antifungal agents, and other nucleotides. PNAs may be used for molecular diagnostics, labelled PNAs are used as hybridization of individuals, e.g. paternity testing, based on hTR gene restriction of individuals, e.g. paternity testing, based on hTR gene restriction fragment length polymorphism (RFLP) pattern. PNAs are also useful as probes to detect the RNA component of a mammalian telomerase and as probes to detect the RNA component of a mammalian telomerase and as inhibitors of telomerase activity. The method of the present invention allows cancerous conditions to be detected with increased confidence and

Example 2; Col 37; 45pp; English.

Query Match 2.9 Best Local Similarity 100. Matches 13, Conservative

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Gaps

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2.9%; Score 13; DB 1; Length 13; 100.0%; Pred. No. 2.8e+02; ive 0; Mismatches 0; Indels

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New peptide nucleic acid (PNA) compounds that inhibit telomerase activity in mammalian cells is useful as probes to detect the RNA component of a mammalian telomerase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Peptide nucleic acid; PNA; telomerase; ribonucleoprotein enzyme; cancer; inhibitor; neoplasia; neurodegenerative disease; aging; hyperplasia; AIDS; HIV; fungal infection; forensic identification; detect; tumour; paternity testing; ss.
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possibly at an earlier stage, before cells are detected as cancerous based on pathological characteristics. The diagnostic and prognostic methods of the present invention can be used to detect an immortal or neoplastic cell or tumour tissue or cancer of any origin, provided the cell expresses telomerase activity and its RNA component.
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                                                                                                                                            2.9%; Score 13; DB 1; Length 13;
100.0%; Pred. No. 2.8e+02;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                        PNA sequence #51 used to inhibit telomerase activity.
                                                                                                            Sequence 13 BP; 1 A; 6 C; 2 G; 4 T; 0 U; 0 Other;
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nes 13; Conservative
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continuation of the continuation with antineoplastic and other cytocatic diseases characterized by abnormal telomere metabolism or telomerase diseases characterized by abnormal telomere metabolism or telomerase cutivity, in combination with antineoplastic and other cytocatic or record of the continuation of continuation can be used to detect an immortal or neoplastic cell or tumour tissue or cancer of any origin, provided the continuation continuation can be used to detect an immortal or continuation continuation can be used to detect an immortal or continuation continuation can be used to detect an immortal or continuation can be used to detect an immortal or continuation can be used to detect an immortal or continuation can be used to detect an immortal or continuation can be continuated to cancer of any origin, provided the continuation cancer of continuation cancer of cancer of cancer 
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immunodeficiency virus (HIV) infection/AIDS (acquired immunodeficiency
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"Phosphorothioate internucleotide linkages"
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100.0%; Pred. No. 2.8e+02;
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les 13; Conservative 0
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control sequence alongside a peptide nucleic acid molecule which hybridises to the mRNA component of mammalian telomerase, and inhibits celomerase activity. Telomerase is a ribonucleoprotein enzyme that synthesizes one strand of the telomeric DNA, using as a template an 11 invention relates to PNA molecules having a sequence of no more than 25 bases, which include the sequence GTTAGG. The uncharged nature of the PNA cackbone increases the melting temperature of associating strands, and affords greater resistance of degradation by proteases or nucleases. The therapeutic PNAs may be used for treating disease conditions such as cancers, neoplasia, hyperplasia, neurodegenerative diseases, aging, human immunodeficiency virus (HIV) infection/AIDS (acquired immunodeficiency cyrolated pathologies, fungal infections, and other cytostatic agents, antifungal agents, and other nucleotides. PNAs may be used for molecular diagnostics, labelled PNAs are used as hybridization cytostatic agents, antifungal agents, and other nucleotides. PNAs may be used for molecular diagnostics, labelled PNAs are used as hybridization of individuals, e.g. paternity testing, based on hTR gene restriction of individuals, e.g. paternity testing, based on hTR gene restriction fragment length polymorphism (RRLP) pattern. PNAs are also useful as probes to detect the RNA component of a mammalian telomerase and as allows cancerned cancerned and cancerned cancerne
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/*tag= = "Peptide nucleic acid molecule, where N-(2-
mainoethyl) glycine units are linked to nuclectide bases
via glycine amino N through a methylenecarbonyl linker"
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AAA37597 standard; DNA; 13 BP.
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The present sequence represents a peptide nucleic acid molecule which hybridises to the mRNA component of mammalian telomerase, and inhibits call hybridises to the mRNA component of mammalian telomerase, and inhibits call the care the telomerase, and inhibits are telomerases activity. Telomerase is a ribonucleoprotein enzyme that synthesizes one strand of the telomeric DNA, using as a template an 11 nucleotide sequence contained within the RNA component of the enzyme. The invention relates to PNA molecules having a sequence of no moore than 25 bases, which include the sequence GTTAGG. The uncharged nature of the PNA careckbone increases the melting temperature of associating strands, increases the rate of association with targeted nucleic acids, and affords greater resistance of degradation by proteases or nucleases. The charapeutic PNAs may be used for treating disease conditions such as cancers, neoplasia, hyperplasia, neurodegenerative diseases or nucleases. The charapeutic PNAs may be used for infection/AIDS (acquired immunodeficiency virus (HIV) antineoplastic and other cytcoxic or cytostatic agents, antifungal agents, and other nucleotides. PNAs may be used for molecular diagnostics, labelled PNAs are used as hybridization of individuals, e.g. paternity testing, based on HTR gene restriction of individuals, e.g. paternity testing, based on HTR gene restriction of individuals, and component of a mammalian telomerase and as through the component of a mammalian telomerase and and probes to detect the RNA component of a mammalian telomerase and and noted and an allows cancerous conditions to be detected with increased conditions and an analysis of the present invention and allows cancerous conditions to be detected with increases and as an angent and an analysis and an areal and as an angent and a
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                                                                                                                                                                       (PNA) compounds that inhibit telomerase activity
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                                                                                                                                                                                              in mammalian cells is useful as probes to detect the RNA component of a mammalian telomerase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             possibly at an earlier stage, before cells are detected as cancerous based on pathological characteristics. The diagnostic and prognostic methods of the present invention can be used to detect an immortal or neoplastic cell or tumour tissue or cancer of any origin, provided the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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                                                                          Corey DR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cell expresses telomerase activity and its RNA component
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PNA sequence #12 used to inhibit telomerase activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 13 BP; 2 A; 5 C; 1 G; 5 T; 0 U; 0 Other;
                                                                          Norton JC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2.9%; Score 13; DB 1; L
.00.0%; Pred. No. 2.8e+02;
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                                                                       Shay JW,
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                                                                                                                                                                                                                                                                      Example 2; Col 37; 45pp; English.
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                                                                          Piatyszek MA,
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                         (TEXA ) UNIV TEXAS SYSTEM
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Best Local Similarity 100.
Matches 13; Conservative
                                                                                                                                                                         New peptide nucleic acid
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                                                                                                                        WPI; 2000-292432/25.
                                                                          Wright WE,
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Peptide nucleic acid; PNA; telomerase; ribonucleoprotein enzyme; cancer;

PNA sequence #4 used to inhibit telomerase activity.

(first entry)

15-AUG-2000

inhibitor, neoplasia, neurodegenerative disease, aging, hyperplasia, AIDS, HIV, fungal infection, forensic identification, detect, tumour,

paternity testing; ss.

Synthetic.

/*tag= a //note= "Peptide nucleic acid molecule, where N-(2-aminoethyl)glycine units are linked to nucleotide bases via glycine amino N through a methylenecarbonyl linker"

Location/Qualifiers

misc_feature

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AAA37547 standard; DNA; 13 BP.
44 GICTAACCCIAAC 56
            GICTAACCCTAAC 1
                   13
                                                                                      AAA37547;
                                                RESULT 374
                                                         AAA37547/c
ID AAA37
XX
AC AAA37
 à
                  g
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DB 1; Length 13;

Score 13;

Query Match

cell expresses telomerase activity and its RNA component

Seguence 13 BP; 5 A; 1 C; 4 G; 3 T; 0 U; 0 Other;

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The present sequence represents a peptide nucleic acid molecule which hybridises to the mRNA component of mammalian telomerase, and inhibits telomerase activity. Telomerase is a ribonucleoprotein enzyme that synthesizes one strand of the telomeric DNA, using as a template an 11 nucleotide sequence contained within the RNA component of the enzyme. The incloude the pNA manying a sequence of no more than 25 backbone increases the melting temperature of associating strands, increases the rate of association with targeted nucleic acids, and affords greater resistance of degradation by proteases or nucleases. The increases the melting temperature of associating strands, increases the rate of association with targeted nucleic acids, and affords greater resistance of degradation by proteases or nucleases. The increases the rate of association with the argeted nucleic acids, and affords greater resistance of degradation by proteases or nucleases. The increases that are target of a canceria, mental acid manucleficiency virus (HTV) infection/ALDS (acquired immunodeficiency syndrome) and associated pathologies, fungal infections, and other diseases, aging, human cativity, in combination with antinooplastic and other cytocoxic or cytostatic agents, antifungal agents, and other nucleotides. PNAs may be cytostatic agents, antifungal agents, and other nucleotides. PNAs may be used for molecular diagnostics, labelled PNAs are used as hybridization probes to detect to quantitate polynucleotides having a human telomerase and individuals, e.g. paternity testing, based on hrm gene restriction frament length polymorphism (RPLP) pattern. PNAs are also useful as probes to detect the RNA component of a mammalian telomerase and as inhibitors of telomerase activity. The method of the present invention can be used to detected as cancerous conditions to be detected with increased confidence and nephastic cell or truncur tissue or cancer of any origin, provided the conditions are also used to detect an immortal or neoplastic cell expresses 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                v peptide nucleic acid (PNA) compounds that inhibit telomerase activity mammalian cells is useful as probes to detect the RNA component of a
                                          Corey DR;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 6; Col 71; 45pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     97US-00838545.
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misc_feature
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Corey DR;

97US-00838545. 96US-00630019.

09-APR-1997; 09-APR-1996;

JS6046307-A 04-APR-2000 (TEXA) UNIV TEXAS SYSTEM.

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Score 13; DB 1; Length 13;
Pred. No. 2.8e+02;
                 100.0%; Pred. No. 2.8
ive 0; Mismatches
   2.9%;
                   Local Similarity 100.
nes 13; Conservative
   Query Match
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The present sequence represents a peptide nucleic acid molecule which hybridises to the mRNA component of mammalian telomeraes, and inhibits telomerase activity. Telomerase is a ribonucleoprotein enzyme that telomerase scrivity. Telomerase is a ribonucleoprotein enzyme that telomerase activity. Telomerase is a ribonucleoprotein enzyme that telomerase so the telomeric DNA, using as a template an 11 nucleotide sequence contained within the RNA component of the enzyme. The invention relates to PNA molecules having a sequence of no more than 25 backbone increases the melting temperature of associating strands, increases the rate of association with targered nucleic acids, and a flords greater resistance of degradation by proteases or nucleases. The therapeutic PNAs may be used for treating disease conditions such as therapeutic PNAs may be used for treating disease conditions such as cancers, neoplasia, hyperplasia, neurodegenerative diseases, aging, human diseases characterized pathologies, fungal infections, and other diseases characterized by abnormal telomere metabolism or telomerase activity, in combination with antineoplastic and other cytotoxic or cycostatic agents, antifungal agents, and other nucleotides. PNAs may be used for molecular diagnostics, labelled PNAs are used as hybridization
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New peptide nucleic acid (PNA) compounds that inhibit telomerase activity in mammalian cells is useful as probes to detect the RNA component of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           probes to detect or quantitate, polymucleotides having a human telemerase RNA (hTR) sequence. PNA probes are also used for forensic identification of individuals, e.g. paternity testing, based on hTR gene restriction fragment length polymorphism (RFLP) pattern. PNAs are also useful as probes to detect the RNA component of a mammalian telomerase and as inhibitors of telomerase activity. The method of the present invention allows cancerous conditions to be detected with increased confidence and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                possibly at an earlier stage, before cells are detected as cancerous based on pathological characteristics. The diagnostic and prognostic methods of the present invention can be used to detect an immortal or neoplastic cell or tumour tissue or cancer of any origin, provided the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Shay JW, Norton JC,
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The present invention relates to peptide nucleic acids (PNAs), comprising a sequence of 6-25 nucleobases, that inhibit telomerase activity in mammalian cells by hybridising to the RNA component of mammalian telomerase and as inhibitors of telomerase activity, or to detect and/or quantitate polynucleotide having the human telomerase RNA component (hTR) sequence, as well as in forensic identification of individuals, such as perernity testing or identification of criminal suspects or unknown descendants based on the hTR gene RFLP pattern. The suspects or unknown descendants based on the hTR gene RFLP pattern. The pythoproliferative diseases, and unknown descendants or preventing cancer, inflammation, lymphoproliferative diseases, and commune disease, or neurodegenerative diseases, are prostatic agents) can be used for treating neoplasia, human immunodeficiency virus (HIV) infections, acquired immunodeficiency syndrome (AIDS) and associated pathologies, and other
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/note= "This sequence is a peptide nucleic acid, i.e. it
contains a polyamide backbone instead of a deoxyribose
backbone"
                Gaps
                                                                                                                                                                                                                                                                                                         Mammalian; peptide nucleic acid; probe; forensic; paternity testing; human telomerase RNA component; hTR gena RFIP pattern; cancer; inflammation; lymphoproliferative disease; autoimmune disease; neurodegenerative disease; neoplasia; hyperplasia; HIV; AIDS; human immunodeficiency virus; acquired immunodeficiency syndrome; telomere metabolism; mutant; cytostatic; anti-inflammatory; immunosuppressive; polyamide backbone; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New peptide nucleic acids that hybridizes to the RNA component of mammalian telomerase, useful for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or
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                                                                                                                                                                                                                                                                          PNA 12 inhibiting human and mammalian telomerase activity.
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                Indels
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Pred. No. 2.8e+02;
              Mismatches
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                                                                                                                                                                       AAS15468 standard; DNA; 13 BP.
100.08;
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                13; Conservative
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Best Local Similarity
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Synthetic.
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              Matches
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      activity. The present sequence represents one of the PNA sequences of the invention
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/*tag= a
/note= "This sequence is a peptide nucleic acid, i.e. i
contains a polyamide backbone instead of a deoxyribose
backbone"
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diseases characterised by abnormal telomere metabolism or telomerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New peptide nucleic acids that hybridizes to the RNA component of mammalian telomerase, useful for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or
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                                                                                           Length 13;
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                                                             Sequence 13 BP; 2 A; 5 C; 1 G; 5 T; 0 U; 0 Other;
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                                                                                             DB 1;
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                                                                                           2.9%; Score 13;
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nes 13; Conservative
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Dynamic of the page of treating or preventing cancer, information, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases, an combination with other pharmaceuticals (such as diseases. The PMAs in combination with other pharmaceuticals (such as antineoplastic or cytostatic agents) can be used for treating neoplasia, hyperplasia, human immunodeficiency virus (HIV) infections, acquired immunodeficiency syndrome (AIDS) and associated pathologies, and other diseases characterised by abnormal telomere metabolism or telomerase activity. The present sequence represents one of the PNA sequences of the invention. Note: The present sequence represents SED ID No 4 in the SEQ ID Internal and column 4. However, table 1 in column 29 shows three sequences with SEQ ID No 4. The first SEQ ID No 4 (PNA #6, AASI5433) appears to be identical to SEQ ID No 1 in the claims and the SEQ ID isting. The second SEQ ID No 4 in table 1 appears to be identical to SEQ ID 4 is the present sequence and appears to be identical to SEQ ID 4 in the SEQ ID No 4 in the present sequence and appears to be identical to SEQ ID 4 is the present sequence and appears to be identical to SEQ ID No 4 in the
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/note= "This sequence is a peptide nucleic acid, i.e. it
contains a polyamide backbone instead of a deoxyribose
backbone"
or unknown descendants based on the hTR gene RFLP pattern. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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Pred. No. 2.8e+02;
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100.0%;
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Best Local Similarity 100.c
Local 13; Conservative
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modified base
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09-APR-1997;
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The present invention relates to peptide nucleic acids (PNAs), comprising a sequence of 6-25 nucleobases, that inhibit telomerase activity in nammalian cells by hybridising to the RNA component of mammalian cells by hybridising to the RNA component of mammalian telomerase and as inhibitors of telomerase activity, or to detect and/or quantitate polynucleotide having the human telomerase RNA component (HTR) sequence, as well as in forensic identification of individuals, such as paternity testing or identification of criminal suspects or unknown descendants based on the hTR gene RFLP pattern. The suspects or unknown descendants based on the hTR gene RFLP pattern. The suspects or unknown descendants or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases. The PNAs in combination with other pharmaceuticals (such as antineoplastic or cyclostatic agents) can be used for treating neoplasia, hyperplasia, human immunodeficiency virus (HIV) infections, acquired immunodeficiency syndrome (AIDS) and associated pathologies, and other diseases characterised by abnormal telomera metabolism or telomerase.
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/note= "This sequence is a peptide nucleic acid, i.e. it
contains a polyamide backbone instead of a deoxyribose
backbone"
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                inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases.
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mammalian telomerase, useful for treating or preventing cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PNA 13 inhibiting human and mammalian telomerase activity.
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100.0%; Pred. No. 2.8e+02;
Live 0; Mismatches 0; Indels
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                                                                            Claim 7; Col 73; 46pp; English.
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Best Local Similarity 100.
Matches 13, Conservative
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09-APR-1997;
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/note= "This sequence is a peptide nucleic acid, i.e. it
contains a polyamide backbone instead of a deoxyribose
backbone"
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                                                                                                                New peptide nucleic acids that hybridizes to the RNA component of mammalian telomerase, useful for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or
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                                        Corey DR, Norton JC;
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                                        Piatyszek MA,
                                                                                                                                                                                                                        Example 2; Col 37-38; 46pp; English.
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                                                                                                                                                                               neurodegenerative diseases.
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(TEXA ) UNIV TEXAS SYSTEM.
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                                                                              WPI; 2001-638024/73
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Best Local Similarity
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                                        Shay JW,
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AAS15465/c
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mammalian cells by hybridising to the RNA component of mammalian cells by hybridising to the RNA component of mammalian cells by hybridising to the RNA component of mammalian cells by hybridising to the RNA component of mammalian telomerase. The PNAs are useful as probes to detect the RNA component of detect and/or quantitate polynucleoxide having the human telomerase RNA component (hTR) sequence, as well as in forensic identification of criminal cuspects or unknown descendants based on the hTR gene RFLP pattern. The PNA can be further used for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases. The PNAs in combination with other pharmaceuticals (such as antineoplastic or cytostatic agents) can be used for treating neoplasia, human immunodeficiency virus (HIV) infections, acquired immunodeficiency syndrome (ADS) and associated pathologies, and other diseases characterised by abnormal telomerame etabolism or telomerase activity. The present sequence represents one of the PNA sequences of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to peptide nucleic acids (PNAs), comprising
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New peptide nucleic acids that hybridizes to the RNA component of mammalian telomerase, useful for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases.
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                                                                                                                                                                                                                                                                                                                                   Norton JC;
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97US-00838545
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Matches 13; Conservative
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09-APR-1997;
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25-SEP-2001
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                                                                                                                                                                                                                                                                                                                                   Shay JW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 380
AAS15423/c
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US09436060A-16.rng.sl

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The present invention relates to peptide nucleic acids (FNAs), comprising a sequence of 6-25 nucleobases, that inhibit telomerase activity in cammalian cells by hybridising to the RNA component of mammalian telomerase and as inhibitors of telomerase activity or to telomerase. The PNAs are useful as probes to detect the RNA component of mammalian telomerase and as inhibitors of telomerase activity, or to detect and/or quantitate polynucleotide having the human telomerase RNA component (hTR) sequence, as well as in forensic identification of individuals, such as paternity testing or identification of criminal suspects or unknown descendants based on the hTR gene RFLP pattern. The PNA can be further used for treating or preventing cancer, inflammation, lymphoppicliferative diseases, autoinmune disease, or neurodegenerative diseases. The PNAs in combination with other pharmaceuticals (such as antineophastic or oyfostatic agents) can be used for treating neoplasia, hyperplasia, human immunodeficiency virus (HIV) infections, acquired immunodeficiency syndrome (AIDS) and associated pathologies, and other diseases characterised by abnormal telomere metabolism or telomerase activity. The present sequence represents one of the PNA sequences of the invention. Note: The present sequence represents SED ID No 1 but is shown as the first SEQ ID No 4 in table 1 (column 29)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mammalian, forensic, paternity testing, human telomerase RNA component, hTR gene RFLP pattern; cancer; inflammation; lymphoproliferative disease; autofimmune disease; neurodegenerative disease; neurojamina; hyperplasia; HIV; AlDS; human immunodeficiency virus; telomere metabolism; mutant; acquired immunodeficiency syndrome; cytostatic; anti-inflammatory;
/*tag= a
/*tag= a
/*tag= "This sequence is a peptide nucleic acid, i.e. it
contains a polyamide backbone instead of a deoxyribose
backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Phosphorothioate (PS) oligomer III used to inhibit telomerase activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                             New peptide nucleic acids that hybridizes to the RNA component of mammalian telomerase, useful for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                                                                                                                                                                                                                                                 Shay JW, Wright WE, Piatyszek MA, Corey DR, Norton JC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2.9%; Score 13; DB 1; Length 13;
100.0%; Pred. No. 2.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 13 BP; 3 A; 1 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Prea. ...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 7; Col 73; 46pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAS15459 standard; DNA; 13 BP
                                                                                                                                                                                      99US-00349532.
                                                                                                                                                                                                                           96US-00630019.
97US-00838545.
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                                                                                                                                                                                                                                                                                          (TEXA ) UNIV TEXAS SYSTEM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CTAACCCTAACTG 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CTAACCCTAACTG 1
                                                                                                                                                                                                                                                                                                                                                                          WPI; 2001-638024/73.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity
                                                                                                      US6294650-B1
                                                                                                                                                                                      08-JUL-1999;
                                                                                                                                                                                                                               09-APR-1996;
                                                                                                                                                                                                                                                 09-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            14-FEB-2002
                                                                                                                                               25-SEP-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 381
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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The present invention relates to peptide nucleic acids (PNAs), comprising a sequence of 6-25 nucleobases, that inhibit telomerase activity in mammalian cells by hybridising to the RNA component of mammalian telomerase and as inhibitors of telomerase activity or to detect and/or quantitate polynucleotide having the human telomerase RNA component (HTR) sequence, as well as in forensic identification of component (HTR) sequence, as well as in forensic identification of individuals, such as paternity testing or identification of runknown descendants based on the hTR gene RFLP pattern. The PNAs an be further used for treating or identification of criminal such as the PNAs in combination with other pharmaceuticals (such as diseases. The PNAs in combination with other pharmaceuticals (such as intimoplastic or cycostatic agents) can be used for treating neoplasis, hymmal immunodeficiency virus (HIV) infections, acquired immunodeficiency syndrome (AIDS) and associated pathologies, and other diseases characterised by abnormal telomerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                               1. .13
//tag=
/label= OTHER
/note= "Phosphorothioate internucleotide linkages"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New peptide nucleic acids that hybridizes to the RNA component of mammalian telomerase, useful for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      activity. The present sequence represents a phosphorothioate (PS) oligomer used to inhibit telomerase activity in the methods of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PNA 2 inhibiting human and mammalian telomerase activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 13 BP; 5 A; 1 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Wright WE, Piatyszek MA, Corey DR,
immunosuppressive; phosphorothioate; ss
                                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1; Col 29; 46pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           96US-00630019.
97US-00838545.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             99US-00349532.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (TEXA ) UNIV TEXAS SYSTEM.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              42 TIGICIAACCCTA 54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity 100.
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2001-638024/73
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                                                                                                                                                                   Key
modified_base
                                                                    sapiens
                                                                                                                                                                                                                                                                                                                                                                  US6294650-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             08-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              09-APR-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             09-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             14-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                            25-SEP-2001.
                                                                                                      Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Shay JW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAS15464;
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ID AAS154
XX
AC AAS154
XX
XX
DT 14-FEE
XX
DE PNA 2
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                                                                       Homo
      SON COURT OF COURT OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ð
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2'-arabino-fluorooligonucleotide; phosphoramidate; telomerase; inhibitor; infection; cancer; diagnosis; therapy; cytostatic; virucide; antisense;

/mod_base= OTHER /note= "2'-arabino-fluoronucleosides"

Location/Qualifiers

2. .13 /*tag= a

modified base

Key

antigene; ss.

Synthetic

/mod_base= OTHER /note= "phosphoramidate linkage"

2. .13 /*tag= b

modified base

19-JAN-2001; 2001WO-US001918. 21-JAN-2000; 2000US-0178248P.

WO200153307-A1.

26-JUL-2001

Gryaznov S, Schultz RG; WPI; 2001-589652/66.

(GERO-) GERON CORP.

Phosphoramidate-linked 2'-arabino-fluorooligonucleotide.

(first entry)

26-NOV-2001

AAH26730;

AAH26730 standard; DNA; 13

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The present invention relates to peptide nucleic acids (PNAs), comprising a sequence of 6-25 nucleobases, that inhibit telomerase activity in mammalian cells by hybridising to the RNA component of mammalian telomerase. The PNAs are useful as probes to detect the RNA component of mammalian telomerase and as inhibitors of telomerase activity, or to detect and/or quantitate polynucleotide having the human telomerase RNA component (hTR) sequence, as well as in forensic identification of individuals, such as paternity testing or identification of criminal suspects or unknown descendants based on the NTR gene RFLP pattern. The PNA can be further used for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases. The PNAs in combination with other pharmaceuticals (such as attended human immunodeficiency virus (HIV) infections, acquired immunodeficiency syndrome (AIDS) and associated pathologies, and other diseases characterised by abnormal telomere metabolism or telomerase of the activity. The present sequence represents one of the PNA sequences of the
                                                                                                                                                                                                                                                                              /*tag= a
//note= "This sequence is a peptide nucleic acid, i.e. it
contains a polyamide backbone instead of a deoxyribose
backbone"
              Mammalian; peptide nucleic acid; probe; forensic; paternity testing; human telomeraes RNA component; hTR gene RFEP pattern; cancer; inflammation; lymphoproliferative disease; autoimmune disease; neurodegenerative disease; neoplasia; hyperplasia; HIV; AIDS; human immunodeficiency virus, acquired immunodeficiency syndrome; telomere metabolism; mutant; cytostatic; auti-inflammatory; immunosuppressive; polyamide backbone; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New peptide nucleic acids that hybridizes to the RNA component of mammalian telomerase, useful for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Norton JC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 13 BP; 8 A; 1 C; 2 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Corey DR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Wright WE, Piatyszek MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 2; Col 37-38; 46pp; English.
                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 96US-00630019.
97US-00838545.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            99US-00349532
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        neurodegenerative diseases.
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                                                                                                                                                                                                                                                                            *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-638024/73
                                                                                                                                                                                                                                                          modified base
                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 09-APR-1996;
09-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                             US6294650-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                            08-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                  25-SEP-2001
                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Shay Jw,
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Polynucleotides, used to detect and isolate nucleic acids, inhibit function of RNA and telomerase enzymes and to treat e.g. viral infections, contain 2'-arabino-fluoronucleoside(8) linked to

nucleoside(s).

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The present sequence is that of a N3'-P5' 2'-arabino-fluoro phosphoramidate oligonucleotide that is complementary to telomerase RNA. The Oligonucleotide was used to assess the relative efficacy of movel 2'-arabino-fluoro phosphoramidate oligonucleotides and their 2'-ribo fluorooligonucleotide counterparts (see AAH26728-35) for the inhibition of telomerase activity. Novel phosphoramidate 2'-arabino-fluoroligonucleotides are generally more acid stable, more resistant to cellular proteases, and also show greater telomerase inhibition activity than 2'-ribose-fluoro phosphoramidates. They are therefore useful for treating cancer (claimed) and other diseases in which telomerase activity is present at abnormal levels, such as hyperproliferative or autoimmune diseases e.g. psoriasis, rheumacoid arthritis, immune system disordere requiring immunosuppression, and in the treatment of viral infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2.9%; Score 13; DB 1; Length 13;
100.0%; Pred. No. 2.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 13 BP; 5 A; 1 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 6; Page 46; 61pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.08;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13 rrercraacccra 1
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3.3
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0
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100.0%; Pred. No. 2.8e+02;
ive 0; Mismatches 0; Indels
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Gaps

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RESULT 384 AAH26734/C

38 TTTTTTGTCTAAC 50 13; Conservative

Best Local Similarity Matches 13; Conserv

TTTTTGTCTAAC 1

13

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RESULT 383 AAH26730/c

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New polynucleotide useful for inhibiting telomerase activity in cells, of for treating telomerase-mediated condition or disease, such as cancers, tumors, Hodgkin's disease, or inflammatory conditions.
                                                                                                                                                Human; telomerase; hTR; cytostatic; anti-inflammatory; adenocarcinoma; breast; prostate; colon; mixed cell leukaemia; Hodgkin's disease; fertility; inflammatory condition; tumour; cancer; veterinary; immunosuppression; telomerase inhibitor; ss.
                                                                                                               Human telomerase polynucleotide inhibitor #18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 3; Page 32; 48pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30-MAR-2001; 2001WO-US010476.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                31-MAR-2000; 2000US-00540119
AAS15937 standard; DNA; 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gryaznov SM, Pruzan R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-656955/75
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                                                                                                                                                                                                                                                                                                                                                                                                                                WO200174136-A2
                                                                                                                                                                                                                                                                                                              Key
modified_base
                                                                                                                                                                                                                                                      sapiens.
                                                                           27-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     11-OCT-2001,
                                                                                                                                                                                                                                                                          Synthetic.
oligonucleotide that is complementary to telomerase RNA. The arabino-leotide was used to assess the relative efficacy of novel 2'-arabino-fluorophosphoramidate oligonucleotides and their 2'-ribose fluorooligonucleotide counterparts (see AAH26728-35) for the inhibition of telomerase activity. Novel phosphoramidate 2'-arabino-fluorophoramidate activity are generally more acid stable, more resistant to cellular proteases, and also show greater telomerase inhibition activity than 2'-ribose-fluoro phosphoramidates. They are therefore useful for treating cancer (claimed) and other diseases in which telomerase activity is present at abnormal levels, such as hyperproliferative or autoimmune diseases e.g. psoriaeis, rheumacioid arthritis, immune system disorders requiring immunosuppression, and in the treatment of viral infection
                                                                                                                                            2'-ribose-fluorooligonucleotide; phosphoramidate; telomerase; inhibitor; infection; cancer; diagnosis; therapy; cytostatic; virucide; antisense; antigene; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Polynucleotides, used to detect and isolate nucleic acids, inhibit function of RNA and telomerase enzymes and to treat e.g. viral infections, contain 2'-arabino-fluoronucleoside(s) linked to nucleoside(s)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present sequence is that of a 2'-ribose-fluoro phosphoramidate
                                                                                                               Phosphoramidate-linked 2'-ribose-fluorooligonucleotide.
                                                                                                                                                                                                                                                                                                                               /mod_base= OTHER
/note= "2'-ribose-fluoronucleosides"
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                                                                                                                                                                                                                                                                                                                                                                                                                                /note= "phosphoramidate linkage"
                                                                                                                                                                                                                                                                          Location/Qualifiers
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ВР
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AAH26734 standard; DNA; 13
                                                                             (first entry)
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/*tag=
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                                                                             26-NOV-2001
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                                                                                                                                                                                                                                   Synthetic
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/mod_base= OTHER /note= "N3'-P5' phosphoramidate linkages"

Location/Qualifiers

(first entry)

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1. .13 /*tag=

or

Weinrich SL;

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The invention relates to polynucleotide inhibitors (I) and methods for inhibiting telomerase activity. (I) are useful in inhibiting telomerase activity and proliferation of a telomerase positive cell, and in manufacturing a medicament for inhibiting telomerase activity in a monitation of breast, prostate or colon, mixed cell leukamia and in treating telomerase-mediated condition or disease, such as adenocarcinoma of breast, prostate or colon, mixed cell leukamia, useful in treating tumour or in manufacturing a medicament for the treatment of tumour or in manufacturing a medicament for the diagnostic assays for detecting RNA or DNA. Inhibition of telomerase methods of treating cancer and other disorders involving inappropriate expression of telomerase, and in treating veterinary proliferative diseases. Inhibition of telomerase in haematopoietic stem cells is useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     branches of the immune system. The present sequence represents human telomerase polynucleotide inhibitor #18, as described in the method of
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Best Local Similarity
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Gaps

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0; Indels

2.9%; Score 13; DB 1; Length 13; 00.0%; Pred. No. 2.8e+02;

100.0%; Pred. nv.

Best Local Similarity

Matches

Query Match

TTGTCTAACCCTA 54 13; Conservative

42 13

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RESULT 385 AAS15937/c

BP

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New polynucleotide useful for inhibiting telomerase activity in cella, or for treating telomerase-mediated condition or disease, such as cancers, tumors, Hodgkin's disease, or inflammatory conditions.
                                                                                                                                                 Human; telomerase; hTR; cytostatic; anti-inflammatory; adenocarcinoma;
breast; prostate; oclon; mixed cell leukaemia; Hodgkin's disease;
fertility; inflammatory condition; tumour; cancer; veterinary;
immunosuppression; telomerase inhibitor; ss.
                                                                                                                      Human telomerase polynucleotide inhibitor #7.
                                                                                                                                                                                                                                                                                                                                                                                                                   30-MAR-2001; 2001WO-US010476.
                                                                                                                                                                                                                                                                                                                                                                                                                                                31-MAR-2000; 2000US-00540119.
                      926/c
AAS15926 standard; DNA; 13
                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gryaznov SM, Pruzan R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2001-656955/75.
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                                                                                                                                                                                                                                                                                 modified_base
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                                                                                                                                                                                                                                         Synthetic.
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                                                              AAS15926;
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                   AAS15926,
                                                The invention relates to polynucleotide inhibitors (I) and methods for inhibiting telomerase activity. (I) are useful in inhibiting telomerase activity and proliferation of a telomerase positive cell, and in manufacturing a medicament for inhibiting telomerase activity in a cell and in treating telomerase-mediated condition or disease, such as and in treating telomerase-mediated condition or disease, such as and in treating telomerase mediated condition or disease, such as Hodgkin's disease, fertility and inflammatory conditions. (I) are also useful in treating a tumour or in manufacturing a medicament for the treatment of tumour. The polynucleotide inhibitors may also be used in treatment of tumour. The polynucleotide inhibitors and therapeutic methods of treating cancer and other disorders involving inappropriate contivity in cells in vivo is useful in prophylactic and therapeutic methods of treating cancer and other disorders involving inappropriate diseases. Inhibition of telomerase in haematopoletic stem cells is useful for thmunosuppression and for selectively down-regulating specific.
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telomerase polynucleotide inhibitor #2, as described in the method of the
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                                                                                                                             Human; telomerase; hTR; cytostatic; anti-inflammatory; adenocarcinoma; breast; prostate; colon; mixed cell leukaemia; Hodgkin's disease; fertility; inflammatory condition; tumour; cancer; veterinary; immunosuppression; telomerase inhibitor; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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/mod_base= OTHER
/note= "N3'-P5' phosphoramidate linkages"
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100.0%; Pred. No. 2.8e+02;
ive 0; Mismatches 0;
                                                                                                     Human telomerase polynucleotide inhibitor #2.
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Claim 8; Page 36; 48pp; English
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Les 13; Conservative
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Matches
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Weinrich SL;

/note= "N3'-P5' phosphoramidate linkages"

base= OTHER

/*tag= a

Location/Qualifiers

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The invention relates to polynucleotide inhibitors (I) and methods for inhibiting telomerase activity. (I) are useful in inhibiting telomerase activity. (I) are useful in inhibiting telomerase activity and proliferation of a telomerase positive cell, and in manufacturing a medicament for inhibiting telomerase activity in a cell and in treating telomerase. For inhibiting telomerase activity in a cell adenocarcinoma of breast, prostate or colon, mixed cell leukaemia, Hodgkini a disease, fertility and inflammatory conditions. (I) are also useful in treating a tumour or in manufacturing a medicament for the treatment of tumour. The polynucleotide inhibitors may also be used in diagnostic assays for detecting RNA or DNA. Inhibition of telomerase activity in cells in vivo is useful in prophylactic and therapeutic methods of treating cancer and other disorders involving inappropriate expression of telomerase, and in treating veterinary proliferative expression of telomerase in inhibition of telomerase in inhibition of telomerase in inhibitor sequence represents human telomerase polynucleotide inhibitor #7, as described in the method of the
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100.0%; Pred. No. 2.8e+02;
/ative 0; Mismatches 0; Indels
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Claim 8; Page 36; 48pp; English.
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Best Local Simi.
Matches 13;
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Human telomerase polynucleotide inhibitor #16.

(first entry)

27-FEB-2002

AAS15935;

BP.

AAS15935 standard; DNA; 13

AAS15935/c

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The invention relates to polynucleotide inhibitors (I) and methods for inhibiting telomerase activity. (I) are useful in inhibiting telomerase activity. (I) are useful in inhibiting telomerase activity and proliferation of a telomerase positive cell, and in manufacturing a medicament for inhibiting telomerase activity in a cell and in treating telomerase-mediated condition or disease, such as adenocarcinoma of breast, prostate or colon, mixed cell leukaemia. (I) adenocarcinoma of breast, prostate or colon, mixed cell leukaemia. (I) adenocarcinoma of breast, prostate or colon, mixed cell leukaemia. (I) are also useful in treating a tumour or in manufacturing a medicament for the treatment of tumour. The polynucleotide inhibition of telomerase chapters are and therefore in thibition of telomerase contivity in cells in vivo is useful in prophylactic and therapeutic methods of treating cancer and other disorders involving inappropriate expression of telomerase, and in treating veterinary proliferative consenses on the inmune system. The present sequence represents human branches of the immune system. The present sequence represents human
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                                                                                                                                                                                                                                                                                                                                                                                   Human; telomerase; hTR; cytostatic; anti-inflammatory; adenocarcinoma; breast; prostate; colon; mixed cell leukaemia; Hodgkin's disease; fertility; inflammatory condition; tumour; cancer; veterinary; immunosuppression; telomerase inhibitor; ss.
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                                                                           AAS15930 standard; DNA; 13 BP.
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/*tag= a
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modified_base
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RESULT 388
AAS15930/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New polynuclectide useful for inhibiting telomerase activity in cells, for treating telomerase-mediated condition or disease, such as cancers, tumors, Hodgkin's disease, or inflammatory conditions.
                                                                                                                       Human; telomerase; hTR; cytostatic; anti-inflammatory; adenocarcinoma; breast; prostate; colon; mixed cell leukaemia; Hodgkin's disease; fertility; inflammatory condition; tumour; cancer; veterinary; immunosuppression; telomerase inhibitor; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         branches of the immune system. The present sequence represents human telomerase polynucleotide inhibitor #16, as described in the method of
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/*tag= a
/mod_base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gryaznov SM, Pruzan R,
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2.9%; Score 13; DB 1; Length 13; 00.0%; Pred. No. 2.8e+02;

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13; Conservative

Matches

Local Similarity

Query Match

13 AAACAAAAATGT

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The invention relates to polynucleotide inhibitors (I) and methods for inhibiting telomerase activity. (I) are useful in inhibiting telomerase activity of proliferation of a telomerase positive cell, and in manufacturing a medicament for inhibiting telomerase activity in a call and in treating telomerase-mediated condition or disease, such as adenocarcinoma of breast, prostate or colon, mixed cell leukaemia. Hodgkin's disease, fertility and inflammatory conditions. (I) are also useful in treating a tumour or in manufacturing a medicament for the treatment of tumour. The polynucleotide inhibitors may also be used in diagnostic assays for detecting RNA or DNA. Inhibition of telomerase activity in cells in vivo is useful in prophylactic and therapeutic methods of treating cancer and other disorders involving inappropriate expression of telomerase, and in treating veterinary proliferative diseases. Inhibition of telomerase in haematopoietic stem cells is useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     for immunosuppression and for selectively down-regulating specific branches of the immune system. The present sequence represents human telomerase polynucleotide inhibitor #3, as described in the method of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New polynucleotide useful for inhibiting telomerase activity in cells, for treating telomerase-mediated condition or disease, such as cancers, tumors, Hodgkin's disease, or inflammatory conditions.
                                                                                                                                                                                  Human; telomerase; hTR; cytostatic; anti-inflammatory; adenocarcinoma;
breast; prostate; colon; mixed cell leukaemia; Hodgkin's disease;
fertility; inflammatory condition; tumour; cancer; veterinary;
immunosuppression; telomerase inhibitor; ss.
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                                                                                                                                                  Human telomerase polynucleotide inhibitor #3
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AAS15922 Btandard; DNA; 13 BP.
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                                                                                                                                                                                                                                                                                                                                                            modified base
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                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                             27-FEB-2002
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                                                                                                                                                                                                                                                                                                     Synthetic
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                                                                        AAS15922;
RESULT 390
                   AAS15922
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The invention relates to polynucleotide inhibitors (I) and methods for inhibiting telomerase activity. (I) are useful in inhibiting telomerase activity. (I) are useful in inhibiting telomerase activity and proliferation of a telomerase positive call, and in manufacturing a medicament for inhibiting telomerase activity in a cell and in treating telomerase.mediated condition or disease, such as adenocarcinoma of breast, prostate or colon, mixed cell leukaemia, Hodgkini's disease, fertility and inflammatory conditions. (I) are also useful in treating a tumour or in manufacturing a medicament for the treatment of tumour. The polynucleotide inhibitors may also be used in treatment of tumour. The polynucleotide inhibitors of telomerase activity in cells in vivo is useful in prophylactic and therapeutic methods of treating cancer and other disorders involving inappropriate expression of telomerase, and in treating veterinary proliferative confiseases. Inhibition of telomerase in haematopolitic stem cells is useful for immunosuppression and for selectively down-regulating specific confinences of the immune system. The present sequence represents human telomerase polynucleotide inhibitor #4, as described in the method of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New polynucleotide useful for inhibiting telomerase activity in cells, or for treating telomerase-mediated condition or disease, such as cancers, tumors, Hodgkin's disease, or inflammatory conditions.
                                                                                                                                                                                                                                                                       Human; telomerase; hTR; cytostatic; anti-inflammatory; adenocarcinoma; breast; prostate; colon; mixed cell leukaemia; Hodgkin's disease; fertility; inflammatory condition; tumour; cancer; veterinary; immunosuppression; telomerase inhibitor; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /mod_base= OTHER
/note= "N3'-P5' phosphoramidate linkages"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 13 BP; 2 A; 2 C; 8 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                   Human telomerase polynucleotide inhibitor #4.
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                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
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                                                                                                                  ВР
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                                                                                                                  AAS15923 standard; DNA; 13
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/*tag= a
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CGCCTTCCACCGT 1
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                         AAS15923;
                                                                              RESULT 391
AAS15923/c
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Query Match
Best Local Similarity 100.0
Matches 13; Conservative

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Best Local Similarity Matches 13; Conserv

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Gaps ö

2.9%; Score 13; DB 1; Length 13; 100.0%; Pred. No. 2.8e+02; tive 0; Mismatches 0; Indels

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138 CTGCCGCCTTCCA 150

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The invention relates to polynucleotide inhibitors (I) and methods for inhibiting telomerase activity. (I) are useful in inhibiting telomerase activity. (I) are useful in inhibiting telomerase activity and proliferation of a telomerase positive cell, and in manufacturing a medicament for inhibiting telomerase activity in a cell and in treating telomerase-mediated condition or disease, such as adenorated in the condition or disease, such as adenorated in the condition of disease, fertility and inflammatory conditions. (I) are also useful in treating a tumour or in manufacturing a medicament for the treatment of tumour. The polynucleotide inhibitors may also be used in diagnostic assays for detecting RNA or DNA. Inhibition of telomerase activity in cells in vivo is useful in prophylactic and therapeutic methods of treating cancer and other disorders involving inappropriate cypression of telomerase, and in treating veterinary proliferative diseases. Inhibition of telomerase in haematopoietic stem cells is useful for immunosuppression and for melectively down-regulating specific.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New polynucleotide useful for inhibiting telomerase activity in cells, for treating telomerase-mediated condition or disease, such as cancers, tumors, Hodgkin's disease, or inflammatory conditions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; telomerase; hTR; cytostatic; anti-inflammatory; adenocarcinoma; breast; prostate; colon; mixed cell leukaemia; Hodgkin's disease; fertility; inflammatory condition; tumour; cancer; veterinary; immunosuppression; telomerase inhibitor; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /mod_base= OTHER
/note= "N3'-P5' phosphoramidate linkages"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human telomerase polynucleotide inhibitor #6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Weinrich SL;
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                                                                                                                                                                                                                                                                          BP
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                                                                                                                                                                                                                                                                              AAS15925 standard; DNA; 13
CCGCCTTCCACCG 153
                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                         CCGCCTTCCACCG 1
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/*tag=
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modified_base
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Synthetic.
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The invention relates to polynucleotide inhibitors (I) and methods for inhibiting telomerase activity. (I) are useful in inhibiting telomerase activity and proliferation of a telomerase positive cell, and in manufacturing a medicament for inhibiting telomerase activity in a cell can in treating telomerase. Medicament for inhibiting telomerase activity in a cell and in treating telomerase. Medicate condition or disease, such as denocarcinoma of breast, prostate or colon, mixed cell leukaemia, cu seful in treating a tumour or in manufacturing a medicament for the treatment of tumour or in manufacturing a medicament for the cuseful in treating a tumour or in manufacturing a medicament for the diagnostic assays for detecting RNA or DNA. Inhibition of telomerase cutivity in cells in vivo is useful in prophylactic and therapeutic methods of treating cancer and other disorders involving inappropriate captisession of telomerase, and in treating veterinary proliferative confinences. Inhibition of telomerase in hamansuppression and for selectively down-regulating specific branches of the immune system. The present sequence represents human continuation and inhibitor #5, as described in the method of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New polynucleotide useful for inhibiting telomerase activity in cells, or for treating telomerase-mediated condition or disease, such as cancers, tumors, Hodgkin's disease, or inflammatory conditions.
                                                                                                                                                                                                                                                     Human; telomerase; hTR; cytostatic; anti-inflammatory; adenocarcinoma; breast; prostate; colon; mixed cell leukaemia; Hodgkin's disease; fertility; inflammatory condition; tumour; cancer; veterinary; immunosuppression; telomerase inhibitor; ss.
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/note= "N3'-P5' phosphoramidate linkages"
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                                                                                                                                                                                                                  Human telomerase polynucleotide inhibitor #5.
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                                                                                                  AAS15924 standard; DNA; 13 BP.
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/*tag=
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                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
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                                                             RESULT 393
                                                                                   AAS15924/
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DB 1; Length 13; 2.8e+02;

2.9%; Score 13; 100.0%; Pred. No.

Query Match Best Local Similarity

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Gaps . 0

Query Match 2.9%; Score 13; DB 1; Length 13; Best Local Similarity 100.0%; Pred. No. 2.8e+02; Matches 13; Conservative 0; Mismatches 0; Indels

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                                                                                                                                                                                                                                                                                                  Human; telomerase; hTR; cytostatic; anti-inflammatory; adenocarcinoma;
breast; prostate; colon; mixed cell leukaemia; Hodgkin's disease;
fertility; inflammatory condition; tumour; cancer; veterinary;
immunosuppression; telomerase inhibitor; ss.
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                                                                                                                                                                                                                                                                   Human telomerase polynucleotide inhibitor #19.
 0; Mismatches
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                                                                                                                                                            AAS15938 standard; DNA; 13 BP.
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                                   139 TGCCGCCTTCCAC 151
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 13; Conservative
                                                       13 TGCCGCCTTCCAC 1
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 Matches
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                                                                                                                         RESULT 39
AAS15938/
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2.9%; Score 13; DB 1; Length 13;

Query Match

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         and anti-gene diagnostic or therapeutic applications and may be used for treating telomerase-mediated conditions or diseases, such as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Thiophosphoramidate oligonucleotide; virucide; cytostatic; immunosuppressive; contraceptive; RNA inhibitor; telomerase inhibitor; antisense therapy; viral infection; cancer; hyperproliferative disorder; autoimmune disorder; se.
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                                                    Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Thiophosphoramidate oligonucleotide, SEQ ID NO: 2.
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100.08; PI
                                                                                                                                                                                                                                                                                                                             AAF81193 standard; DNA; 13 BP
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99US-0160444P.
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                                                                                                                161 TAGAGCAAACAAA 173
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nes 13; Conservative
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                                                                                                                                                                           TAGAGCAAACAAA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200118015-A1.
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19-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 30-MAY-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gryaznov S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
                                                                                                                                                                           13
                                                                                                                                                                                                                                                                                                                                                                                     AAF81193;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                  RESULT 395
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            , Best Loc
                                                                                                                                                                                                                                                                                              AAF81193,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           셤
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US09436060A-16.rng.sl

Morin GB

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New oligonucleotide conjugate useful in the treatment of e.g. cancers are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention relates to oligonucleotide conjugates where the oligonucleotide is covalently linked to an aromatic group. Compounds of the invention are used in medicine for inhibition of telomerase enzyme activity in cells (preferably expressed by tumour cells such as cells from cancer of the skin, central nervous system, retina and circulating tumours (e.g. leukaemia and lymphoma)). The present DNA sequence is an oligonucleotide used in conjugates of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Match phosphorothioate modified 2'-O-methyl RNA G-core oligonucleotide
                                              Medicine; telomerase; tumour; cancer; leukaemia; lymphoma; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     telomere length; aging; hyperproliferative condition; cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 13; DB 1; Length 13;
Pred. No. 2.8e+02;
Oligonucleotide #1 used in conjugates of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 13 BP; 5 A; 1 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Anderson KP, Bennett CF,
                                                                                                                                                                                                                                                                                                                                                                          Tolman RL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 12; Page 37; 45pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ;
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93WO-US009297.
95US-00403888.
99US-00299058.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2.9%; 2
                                                                                                                                                                                                                                    21-MAR-2002; 2002WO-US009138
                                                                                                                                                                                                                                                                           23-MAR-2001; 2001US-0278322P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    02-JAN-2002; 2002US-00038335
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADB68045 standard; RNA; 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   42 TTGTCTAACCCTA 54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 2.9
Best Local Similarity 100.
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                          자,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            13 Trgrcraacccra 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            telomerase inhibitors.
                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-092850/08
                                                                                                                                                                                                                                                                                                                            (GERO-) GERON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US2003096776-A1.
                                                                                                                                          WO200277184-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29-SEP-1992;
29-SEP-1993;
12-JUN-1995;
23-APR-1999;
                                                                                              Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ISIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           04-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hanecak RC,
                                                                                                                                                                                                                                                                                                                                                                          Gryaznov S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22-MAY-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADB68045;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (ISIS-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                G-core
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 398
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADB68045/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present sequence was synthesised in an example illustrating an invention relating to polynucleotides comprising a non-homopolymeric sequence of nucleoside subunits joined by at least one inter-subunit linkage that is a N3'-P5' thiophosphoramidate. The thiophosphoramidate oligonucleotides retain a high RNA binding affinity and exhibit a much higher acid stability. They are useful for detecting a specific sequence in a sample, by forming a hybridisation complex with the sequence. They are useful for inhibiting function of an RNA in a cell (for inhibiting translation of a mRNA or for inhibiting telomerase enzyme in a cell). They are also useful in the preparation of a medicament for treatment of viral infection or cancer. The oligonucleotides are useful for anti-sense and anti-gene diagnostic or therapeutic applications and may be used for treating telomerase-mediated conditions or diseases, such as hyperproliferative and autoimmune disorders, and for contraceptive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
                                                                                                                                                                                                                                  Thiophosphoramidate oligonucleotide, virucide, cytostatic, immunosuppressive, contraceptive, RNA inhibitor, telomerase inhibitor; antisense therapy, viral infection, cancer, hyperproliferative disorder,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel thiophosphoramidate polynucleotide useful for detection of RNA or DNA having a given target sequence, for inhibiting RNA function in a cell, and for treating cancer and viral infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                        Thiophosphoramidate oligonucleotide, SEQ ID NO: 10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 13 BP; 3 A; 1 C; 5 G; 4 T; 0 U; 0 Other;
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0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matray T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 7; Page 47; 68pp; English
                                                   AAF81195 standard; DNA; 13 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ВР
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               08-SEP-2000; 2000WO-US024688
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                                                                                                                                          (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gryaznov S, Pongracz K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                           autoimmune disorder; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-265967/27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (GERO-) GERON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                     WO200118015-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10-SEP-1999;
19-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          24-MAR-2003
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                                                                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                              AAF81195;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 397
                          AAFB1195/C
AAFB1195/C
AACB1195/C
AACB1195/C
AACB1195/C
AACB1197/C

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1D AAD5(XX
AC AAD5(XX
DT 24-M
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Gaps

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0; Indels

Chiang M, Brown-Driver VL;

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99US-00299058,

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23-APR-1999;
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                                                                                                Ecker DJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               selected
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABZ58497,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
                                                                                                                                                                                           The invention relates to a novel chemically modified oligonucleotide having no more than about 27 nucleic acid base units. The oligonucleotide modulates mammalian telomere length. The chemically modified oligonucleotide of the invention may be useful for modulating the telomere length of a mammalian chromosome, inhibiting the division of a mammalian cell or modulating the effects of aging of a mammalian cell. The oligonucleotides may also useful for treating diseases associated with abnormal telomere length such as aging and hyperproliferative conditions including cancer. The current sequence is that of the "match" phosphorotion modified 2'-O-methyl RNA G-core oligonucleotide of the invention.
                                                                           New chemically modified oligonucleotides, useful for modulating telomere length of a mammalian chromosome, inhibiting the division of a malignant mammalian cell, or modulating the effects of aging of a mammalian cell.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                **tag= c
/mod_base= mSc, OTHER
/note= "OTHER = Optionally 5-methylcytosine or linked to
Lys residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /*tag= b
/mod_base= OTHER
/mod_base= OTHER = Optionally phosphorothioate backbone and
2'-O-methyl sugar modification"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          telomere length; aging; hyperproliferative condition; cancer; 88; PNA; peptide nucleic acid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         note = "OTHER = Optionally linked to Gly residue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2.9%; Score 13; DB 1; Length 13; 100.0%; Pred. No. 2.8e+02; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Match 2'-0-methyl oligonucleotide / peptide nucleic acid.
                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 13 BP; 3 A; 1 C; 5 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /*tag= a
/mod_base= OTHER
                                                                                                                                                          Example 5; Page 6; 10pp; English
  Wyatt JR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          046/c
ADB68046 Btandard; DNA; 13 BP.
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93WO-US009297.
95US-00403888.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 100.
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   46 CTAACCCTAACTG 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CTAACCCTAACTG 1
Vickers TA,
                                       WPI; 2003-606442/57.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     29-SEP-1992;
29-SEP-1993;
12-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  02-JAN-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic
  Ecker DJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADB68046;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 399
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046/ADB66046ADB66046ADB66046ADB66046ADB66046ADB66046ADB66046ADB66046ADB66046ADB66046ADB66046ADB66046ADB66046ADB66046ADB66046ADB66046ADB66046ADB66046ADB6604ADB6604ADB6604ADB6604ADB6604ADB6604ADB6604ADB6604ADB6604ADB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 mammalian hair growth comprises applying telomerase inhibitor to
                                                                                                                                                                                             New chemically modified oligonucleotides, useful for modulating telomere length of a mammalian chromosome, inhibiting the division of a malignant mammalian cell, or modulating the effects of aging of a mammalian cell.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present sequence is that of telomerase inhibitor VI, an example a telomerase inhibitor useful in the method of the invention for reducing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gарв
                                                      Brown-Driver VL;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Indels
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                                                         Chiang M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 13 BP; 3 A; 1 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 13; DB 1; Lo
Pred. No. 2.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bennett CF,
                                                A. Wyatt JR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 4; 13pp; English
                                                                                                                                                                                                                                                                                                                  Example 5; Page 6; 10pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          12-JUN-2002; 2002WO-US018702.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                27-JUN-2001; 2001US-00893252
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Styczynski P, Ahluwalia GS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     497/c
ABZ58497 standard; RNA; 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CTAACCCTAACTG 58
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                   RC, Anderson KJ
J, Vickers TA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   relomerase inhibitor VI
(ISIS-) ISIS PHARM INC
                                                                                                                                       WPI; 2003-606442/57.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (GILL ) GILLETTE CO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2003-221439/21.
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Best Local Similarity
Matches 13; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         the invention.
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unwanted mammalian (human) hair growth. A composition comprising the telomerase inhibitor is applied to the skin to reduce hair growth. The telomerase inhibitor acts by reducing telomerase levels in hair follicles, reducing telomerase mRNA expression or by promoting the erosion of telomeric DNA. It can be used to reduce hair growth of a woman with hirsutism or to reduce androgen stimulated hair growth (both claimed). A composition comprising a telomerase inhibitor reduces hair growth by at least 15% (preferably 20%) when tested in the Golden Syrian
                                                                                                                                                                                                                                                 Hamster assay
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Sequence 13 BP; 3 A; 1 C; 5 G; 0 T; 4 U; 0 Other;

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Gaps
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Score 13; DB 1; Length 13;
Pred. No. 2.8e+02;
                   0; Indels
                   0; Mismatches
2.9%; 2
                                       CTAACCCTAACTG 58
                    13; Conservative
                                                        CTAACCCTAACTG 1
 Query Match
Best Local Similarity
                                       46
                                                         13
                    Matches
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Telomerase template region complementary oligonucleotide SEQ ID NO:3. BP ADM46660 standard; DNA; 13 (first entry) cancer; tumour 01-JUL-2004 ADM46660; RESULT 401 ADM46660/

ss; telomerase; hTR; template region; cytostatic; telomerase inhibition;

Synthetic.

WO2004029277-A2

08-APR-2004

23-SEP-2003; 2003WO-US029730.

25-SEP-2002; 2002US-00255535

(GERO-) GERON CORP.

Morin GB; Tolman RL, Pongracz K, Gryaznov S,

WPI; 2004-329894/30

Novel covalent oligonucleotide conjugates comprising nucleobase and oligonucleotide exactly complementary to sequence within template region of human telomerase RNA, linked through linker, useful as medicine for cancer. treating

Disclosure; SEQ ID NO 3; 50pp; English

activity, comprising a nucleobase (A) and oligonucleotide (O) containing activity, comprising a nucleobase (A) and oligonucleotide (O) containing a sequence of 2-11 nucleotides exactly complementary to a sequence within the the terminance of thuman telomerase RRA, linked through a linker (L). A compound of the invention has cytostatic activity. The compound is useful for inhibiting the activity of telomerase enzyme, which involves contacting the telomerase with the compound. The compound of the invention is also useful for inhibiting the proliferation of the cell is which involves contacting the cell with the compound. The cell is treating cancer. The compound is useful for inhibiting or reducing the compound is useful for inhibiting or reducing telomerase activity e.g., tumour cells. The telomerase-positive tumour cells are from cancer of the skin, connective tissue, adipose, breast, lung, stomach, pancreas, ovary, cervix, uterus, kidney, bladder, colon, prostate, central nervous system (CNS), retina and circulating tumours The invention relates to a novel compound having telomerase inhibition

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /mod_base= OTHER
/note= "OTHER = Optionally attached to C3H711CONH group"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a novel method for manufacturing a labelled nucleic acid which comprises introducing an 11C to the transduction region and contacting the nucleic acid fragment and 11C compound, where the nucleic acid fragment is a polynucleic acid or peptide nucleic acid. The method of the invention may be useful for manufacturing a labelled nucleic acid and provides sensitive detection of a target mRNA, such neuroglia myclibril acidic protein (GFAP). The current sequence is that of the labelled nucleic acid/11C-related PNA-antisense oligo 2 of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          by introducing carbon-11 to
nucleic acid fragment and carbon-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
(such as leukaemia and lymphoma). The present sequence represents an oligonucleotide of the invention.
                                                                                                             Gaps
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                                                                                                                                                                                                                                                                                                                                                                        labelled nucleic acid manufacture; 11C; sensitive detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2.9%; Score 13; DB 1; Length 13; 100.0%; Pred. No. 2.8e+02; ive 0; Mismatches 0; Indels
                                                                          2.9%; Score 13; DB 1; Length 13; 100.0%; Pred. No. 2.8e+02; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                          Labelled nucleic acid/11C-related PNA-antisense oligo 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 13 BP; 3 A; 1 C; 5 G; 4 T; 0 U; 0 Other;
                                               Seguence 13 BP; 5 A; 1 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Manufacturing labeled nucleic acid, transduction region, and contacting
                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 2; Fig 7; 17pp; Japanese.
                                                                                                                                                                                                                                                   BP.
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                                                                                                                                                                                                                                                 ADO21607 standard; DNA; 13
                                                                                                                                                                                                                                                                                                              (first entry)
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                                                                                                                                          42 TIGICIAACCCIA 54
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es 13; Conservative
                                                                                                             13; Conservative
                                                                                                                                                                      13 Trercraaccera 1
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                                                                               Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                          PNA-antisense; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            JP2004123542-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                   Unidentified
                                                                                                                                                                                                                                                                                                              15-JUL-2004
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                                                                                                                                                                                                                                                                               AD021607;
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                                                                                                                                                                                                                      RESULT 402
                                                                                                             Matches
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Diagnosing and treating a patient with a solid tumor that produces folate receptor alpha, comprises administering a folate receptor alpha inducer comprising a steroid receptor agent, optionally with a histone deacetylase inhibitor.
                                                                                                                                                                         Example; SEQ ID NO 3; 99pp; English.
                               17-MAR-2003; 2003US-0455705P.
                                                    (MEDI-) MEDICAL COLLEGE OHIO
                                                                                             WPI; 2004-690896/67.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                          Ratnam M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAV93815
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                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a novel method for manufacturing a labelled nucleic acid which comprises introducing an 11C to the transduction region and contacting the nucleic acid fragment and 11C compound, where the nucleic acid fragment is a polynucleic acid or peptide nucleic acid. The method of the invention may be useful for manufacturing a labelled nucleic acid and provides sensitive detection of a target mRNA, such neuroglia myofibril acidic protein (GFAP). The current sequence is that of the labelled nucleic acid/11C-related PNA-antisense oligo 1 of the
                                                                                                                                                                                                                                                                                                      Manufacturing labeled nucleic acid, by introducing carbon-11 to transduction region, and contacting nucleic acid fragment and carbon-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                 abelled nucleic acid manufacture; 11C; sensitive detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2.9%; Score 13; DB 1; Length 13; 100.0%; Pred. No. 2.8e+02; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FR-alpha; G/C rich; solid tumour; folate receptor alpha; steroid receptor; cancer; cytostatic; ds; Spl binding.
                                                                                             Labelled nucleic acid/11C-related PNA-antisense oligo 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 13 BP; 5 A; 1 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human G/C-rich (Sp1 binding) element SEQ ID NO:3.
                                                                                                                                                                                                                                                                                                                                                   Example 2; Fig 7; 17pp; Japanese.
                              ADO21606 standard; DNA; 13 BP
                                                                                                                                                                                                                                                            (NIKL ) JAPAN STEEL WORKS LTD
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                                                                                                                                                                                                                  30-AUG-2002; 2002JP-00254671.
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                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   42 TTGTCTAACCCTA 54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity 100.
13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13 TTGTCTAACCCTA 1
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                                                                                                                              PNA-antisense; se
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                                                                                                                                                                        JP2004123542-A.
                                                                                                                                                  Unidentified
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                                                                      15-JUL-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               invention
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Best Local Si
Matches 13,
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                                                   ADO21606;
                                                                                                                                                                                                                                                                                                                              compound
          RESULT 403
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                    ADO21606/
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The invention relates to a novel method for diagnosing and treating a patient (suspected of) having a solid tumour that produces folate receptor alpha (Rr-alpha), comprising administering a folate receptor receptor alpha (Rr-alpha), comprising administering a folate receptor alpha inducer comprising at least one steroid receptor agent, optionally together with one or more histone deacetylase inhibitors. The folate receptor alpha inducer increases levels of the receptor on plasma membranes of tumour cells and in at least one type of body fluid. Also claimed as part of the invention are: a method for diagnosing or therapeutically treating a patient comprising activating the folate receptor alpha (RR-alpha) gene in the patient using at leat one of an agonist or antagonist of cestrogen receptor (ER), progesterone receptor (PR), androgen receptor (AR) or glucocorticoid receptor (ER); A method of constantially increasing folate receptor (ER), type alpha expression in a tumour using at least one steroid receptor agent in a manner selective to FR-alpha-positive tissues; and finally a method to enhance comprising use of at least one steroid receptor agent that binds to FR-alpha; and a serum marker for diagnostic screening and monitoring of cancers during and after treatment comprising use of at least one steroid receptor. A method of the cancers during and after treatment comprising and monitoring of cancers during and after treatment comprising at esettor activity. The methods are useful in diagnosing and treating a patient with, or suspected of having, at least one solid tumour. The present sequence represents a human G/C-rich (Spl binding)
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target, substrate; catalyst; modulation; expression; Raf gene; delivery;
screening; identification; synthesis; deprotection; purification; cancer;
inflammation; psoriasis; non-hepatic ascites; infection; genetic drift;
restenosis; rheumatoid arthritis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gape
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100.0%; Pred. No. 2.8e+02;
tive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 13 BP; 1 A; 0 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAV93815 standard; RNA; 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18-FEB-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21 GGGAGGGGTGGTG 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GGGAGGGGTGGTG 13
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Les 13, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAV93815;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    element
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The present invention relates to novel 2',5'-oligoadenylic acid analogs and their pharmacologically- acceptable salts. The analogs are stable with superior antitumour and antiviral activity and so are useful in cancer or antiviral therapy e.g. as antisense drugs. The present sequence was used to illustrate the invention.
                                                                                                              /mod_base= OTHER
/note= "2',4'-oxyethylene linkage in the sugar residues
and phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Stable 2',5'-oligoadenylic acid analogs containing natural and modified nucleic acid units as well as unusual phosphate groups with excellent activity particularly antitumor, applicable in cancer or antiviral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Galanin receptor; GALR1; human; single nucleotide polymorphism; SNP; drug discovery; haplotyping; infectious diarrhoea; growth hormone deficiency; allele-specific oligonucleotide; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Galanin receptor gene GALR1 allele-specific oligonucleotide #46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1; Length 14;
3.1e+02;
hes 0; Indels
                                                                                                                                                                                                               /mod_base= OTHER
/note= "G-hydroxyethyl phosphate"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 14 BP; 5 A; 1 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2.9%; Score 13; DB 100.0%; Pred. No. 3.1 tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 111; 220pp; Japanese.
                                                       Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16-APR-2001; 2001WO-US012306.
                                                                                                                                                                                                                                                                                                                                                    19-NOV-2003; 2003WO-JP014748.
                                                                                                                                                                                                                                                                                                                                                                                           19-NOV-2002; 2002JP-00334731
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AAS98334 standard; DNA; 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            42 TIGICTAACCCIA 54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 2.9
Best Local Similarity 100.
Matches 13; Conservative
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                                                                                                                                                                                              /*tag=
                                                                                              '*tag≃
                                                                                                                                                                                                                                                                                                                                                                                                                                (SANY ) SANKYO CO LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Koizumi M, Morita K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2004-460494/43.
                                                                                                                                                                                                                                                                        WO2004046161-A1
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                                                                         modified base
                                                                                                                                                                          modified_base
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                                                                                                                                                                                                                                                                                                               03-JUN-2004
                  Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAS98334;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 407
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a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A method has been developed to the system a random library of mucleic comprises: (a) introducing into the system a random library of mucleic acid catalystes (MAC) having a substrate binding domain (SBD), comprising a random sequence, and a catalystic domain (CD); and (b) identifying NAC in systems where modulation has occurred and/or determining the sequence of at least part of the SBDs in such systems. Nucleic acid molecules with endomuclease activity and catalytic activity, from the present invention, are used to modulate gene expression in plant and mammalian cells and to cleave target nucleic acid, particularly for treating systemic diseases caused by specific RNA, e.g. cancer, inflammation, psoriasis, non-hepatic ascites and infection. They may also be used to detect genetic drift and mutch RNA-cleaving activity that modulate expression of the Raf gene, are used to treat cancer, restenosis, psoriasis or rheumatoid arthritis, or generally any condition associated with the level of c-raf. Introduction of sugar/phosphate modifications increases stability against nuclease and activity. AAV90922 to AAV93877 repersent NACs that can be used in the method, specifically for modulating the expression of a Raf gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Identifying new catalytic nucleic acid that modulates selected processes - especially ribozymes that cleave Raf RNA for treating cancer, restenosis, and also new ribozymes and modified nucleoside triphosphates used as antiviral agents and synthons.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A method has been developed for the identification of a nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cytostatic, virucide, 2', 5'-oligoadenylic acid analog; antitumour; antiviral; cancer; ss.
                                                                                                                                                                                                                                                                                                                                 Bellon L;
Burgin A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2.9%; Score 13; DB 1; Length 14; 34.6%; Pred. No. 3.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Indels
                                                                                                                                                                                                                                                                                                                               Kisich K,
                                                                                                                                                                                                                                                                                                                             nic J, Reynolds M, Kisich K,
Mcswiggen JA, Karpeisky A,
', Beaudry A, Sweedler D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 14 BP; 1 A; 5 C; 6 G; 0 T; 2 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 179; Page 175; 259pp; English.
                                                                                                                                                   970S-0051718P.
970S-0056808P.
970S-0061321P.
970S-0061324P.
970S-0064866P.
                                                                                                                97US-0046059P.
                                                                                                                                                                                                                                                                                                                                 Matulic-Adamic J,
                                                                             98WO-US009249
                                                                                                                                                                                                                                                        97US-0068212P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              84.6%;
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                                                                                                                                                                                                                                                                                               (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          181 AGCTGCTGCCCC 193
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                                                                                                                                                                                                                                                                                                                                 Jarvis T, Matulic-Adamic
Parry T, Beigelman L, M
Thompson J, Workman CT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AGCUGCUGGCCCG 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1999-009494/01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
  WO9850530-A2
                                                                           05-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             26-AUG-2004
                                                                                                                                                       03-JUL-1997
22-AUG-1997
                                                                                                                                                                                                                                                        19-DEC-1997
                                      12-NOV-1998
                                                                                                                                                                                              02-OCT-1997
                                                                                                                                                                                                               02-OCT-1997
05-NOV-1997
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Stephena JC;

Nandabalan K,

Denton RR,

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WPI; 2002-557737/59.

(GENA-) GENAISSANCE PHARM INC

03-JAN-2001; 2001WO-US000092. 03-JAN-2001; 2001WO-US000092.

US09436060A-16.rng.sl

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                                                                                                                                                                                                                                      The invention relates to genotyping human galanin receptor (GALRI) gene of an individual, involving determining for the two copies of the GALRI of an individual, in the indentity of the nucleotide pair at come or more polymorphic sites. The method is useful for determining whether an individual has a haplotype or haplotype pairs defined in the specification. This is useful for improving the efficacy and reliability of several steps in the discovery and development of drugs for treating diseases associated with GALRI activity, e.g., infectious distributions and growth hormone deficiency, to validate GALRI as a candidate agent for treating a specific condition or disease predicted to be associated with GALRI activity, and in the design of clinical trials of candidate drugs for treating a specific condition or disease predicted to be associated with GALRI activity. The method is useful to compounds targeting GALRI activity. The method is useful to compounds the expression and function of GALRI, and in expressing GALRI protein for use in screening for candidate drugs to treat diseases related to GALRI activity. The polymucleotide or variant is useful for studying expression of the polymucleotide or variant is useful for studying expression of the biological activity of GALRI as well as on the binding the affantity of candidate drugs terrepression on the biological activity of GALRI as well as on the binding confining and for studying the effect of the variation on the biological activity of GALRI as well as on the binding caperesent human GALRI gene allele-specific cligonucleotides used to accept the capered of the capered of the method of the capered of the method of the capered of the cape
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                                                                                                                                             Genotyping human galanin receptor gene of an individual for determining haplotype of an individual, involves determining the identity of nucleotide pair at specific polymorphic sites for two copies of the gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Lecithin-cholesterol acyltransferase; LCAT; Norum disease; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        fish-eye disease, atherosclerotic cardiovascular disease, forensic, population diversity, anthropological lineage, paternity testing, human, polymorphism, allele-specific oligonucleotide, ASO; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                               Denton RR, Nandabalan K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 13; DB 1; Length 15;
Pred. No. 3.4e+02;
1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human LCAT gene polymorphism detection ASO primer #16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 15 BP; 2 A; 3 C; 9 G; 0 T; 0 U; 1 Other;
                                                                               Choi JY,
                                                                                                                                                                                                               Claim 16; Page 15; 99pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABK97507 standard; DNA; 15 BP.
                                                (GENA-) GENAISSANCE PHARM INC
                 14-APR-2000; 2000US-0197838P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 2.9%;
Best Local Similarity 86.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      82 TTGCTCCCGCGCGC 96
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       13; Conservative
                                                                               Chew A,
                                                                                                                WPI; 2002-066341/09.
                                                                               Bentivegna SC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200253575-A1
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The present invention relates to a new polynucleotide comprising a nucleotide sequence which is a polymorphic variant of a reference sequence which is a polymorphic variant of a reference is useful for lecithin-cholesterol acyltransferase (LCAT). The invention is useful for identifying an association between a trait (preferably a clinical response to drug targeting LCAT) and at least one genotype or haplotype of LCAT gene. The method of the invention has applicability in developing diagnostic tests and therapeutic treatments for Norum disease. It haplotyping and genotyping methods are useful for studying population if ineage at the phenotypic level, paternity testing, forensic applications and for identifying association between the LCAT genetic applications and for identifying association between the LCAT genetic applications trait such as level of drug response or susceptibility to disease. In caddition, the methods for identifying the LCAT haplotypes present in cindividuals are useful in the development of drugs targeting LCAT. For example, determining the frequency of individual LCAT haplotypes in a copulation with a specific disease, e.g. Norum disease, will facilitate the development of drugs targeting the LCAT isoform(s) that are most trepresent in that disease population. The present nucleic acid sequence trepresents one of a collection (ABK97492-ABK9515) of individual expecience.
Novel isolated polymorphic variant polynucleotide of lecithin-cholesterol acyltransferase gene, useful for studying expression and biological function of the gene, and for therapeutic, diagnostic or forensic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       that were used in the invention to detect
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single nuclectide polymorphism; SNP; drug screening; atherosclerosis;
cardiovascular disease; hypercholesterolaemia;
allele specific oligonuclectide; 88; probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    .9%; Score 13; DB 1; Length 15; 0.0%; Pred. No. 3.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 100.0%; Pred. No. 3.4
Matches 13; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  oligonucleotide (ASO) primers that we polymorphisms in the human LCAT gene
                                                                                                                                                                                                                          Claim 16; Page 17; 72pp; English.
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ID ADG98425 standard; DNA; 15
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Matches
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                                                                                                                                                                                                                             RESULT
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                                                                                                                                                               The invention comprises the amino acid and coding sequences of the human cholesteryl ester transfer protein (CETP), the invention also comprises polymorphisms identified within the CETP gene. The DNA and protein sequences of the invention are useful in haplotyping and/or genotyping the CETP gene in an individual. The DNA and protein sequences may also be used to screen drugs or compounds targeting the CETP or its variant to treat a condition or disease associated with CETP (e.g. atherosclerosis, cardiovascular disease or hypercholesterolaemia). The present DNA sequence represents an allele specific oligomucleotide probe for the
                                                                                                 New isolated polynucleotide useful for haplotyping and/or genotyping cholesteryl ester transfer protein (CETP) gene in an individual or in screening for drugs useful in treating diseases associated with CETP
                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Porcine reproductive and respiratory syndrome virus; PRRSV; vaccine; antigen; polymerase chain reaction; PCR; primer; ss.
                                                         Parks KE;
                                                                                                                                                                                                                                                                                                            ..
                                                                                                                                                                                                                                                                                       Score 13; DB 1; Length 15;
Pred. No. 3.4e+02;
1; Mismatches 1; Indels
                                                         Lee HH,
                                                                                                                                                                                                                                                                   Sequence 15 BP; 4 A; 4 C; 5 G; 1 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Lum MA;
                                                       Lachowicz M,
Tirrell C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Morozov I,
                                                                                                                                               Claim 43; SEQ ID NO 57; 250pp; English.
                                                      Kazemi A, L
Sausker EA,
                                    (GENA-) GENAISSANCE PHARM INC
28-APR-2003; 2003WO-US013288.
                  26-APR-2002; 2002US-0375791P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                PRRSV sequencing primer DP966.
                                                                                                                                                                                                                                                                                       2.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Halbur P,
                                                                                                                                                                                                                                                                                                                           310 GCTCTGTCAGCGGCG 324
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    94US-00301435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    95WO-US010904
                                                                                                                                                                                                                                                                                                                                     15 GCTCTGTYAGCCTCG 1
                                                                                                                                                                                                                                                                                                                                                                                         AAT14404 standard; DNA; 16
                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                        13; Conservative
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                                                      i, Chew Rounds
                                                                                 WPI; 2003-865576/80.
                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
Matches 13; Conserv
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MENG X.
HALBUR P.
MOROZOV I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Paul PS, Meng X,
                                                                                                                                                                                                                                                    human CETP gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LUM M A.
                                                       Anastasio AE,
                                                                Petersen N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO9606619-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-SEP-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                             05-AUG-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                07-MAR-1996.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                              activity
                                                                                                                                                                                                                                                                                                                                                                                                            AAT14404;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (PAUL/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (MENG/)
(HALB/)
(MORO/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (LUMM/)
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                                                                                                                                                                       Primer DP966 (AAT14404) is specific to porcine reproductive and respiratory syndrome virus (PRRSV). It was used with other sequencing primers (AAT1431-182) to determine the sequences of the putative membrane (M) and nucleocapsid (N) genes of PRRSV isolate ISU-12 (see also AAT14391-92) and of 5 other American PRRSV isolates (AAT14405-09) and Buropean
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
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New porcine reproductive and respiratory syndrome virus DNA - and proteins encoded by open reading frames of an Iowa strain of the virus; are used in vaccines against PRRSV in pigs.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Enzymatic and antisense nucleic acid inhibition of repressor genes, useful for producing e.g. granulocyte colony stimulating factor protein, interferon alpha and erythropoietin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 2.9%; Score 13; DB 1; Length 17; Best Local Similarity 100.0%; Pred. No. 3.98+02; Matches 13; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                           Sequence 16 BP; 2 A; 4 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 17 BP; 5 A; 9 C; 2 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    2.9%; Score 13; DB .00.0%; Pred. No. 3.6 .ve 0; Mismatches
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                                                                                                                        Disclosure; Page 77; 228pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 37; Page 56; 164pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hammerhead ribozyme substrate #4.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.08;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99US-0129390P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAF01709 standard, DNA; 17
                                                                                                                                                                                                                                                                                                                                     strain Lelystad (AAT14410)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            268 GGGCTTCTCCGG 280
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4 GGGCTTCTCCGG 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2000-647423/62.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            interferon alpha
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The sequences given in AAT80211-382 represent synthetic oligonucleotides which are complementary to a portion of the 5' untranslated region (UTR) of hepatitis C virus (HCV). These sequences may be used in a pharmaceutical composition for the control or prevention of HCV infection. They may be used to inhibit replication or expression of HCV in a sample. They may be used to inhibit HCV in the control or expression of HCV in thibit HCV replication in a cell and are therefor useful in the treatment of HCV infections such as chronic and acute hepatitis and hepatocarcinoma. This sequence forms multiplex binding complexes with regions of the HCV genome. This sequence forms a duplex at the region - 227 and forms a purine strand triplex at the region - 212 to -222
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            go:nucleotide(s) complementary to HCV 5' untranslated region - used intreatment and detection of HCV infection, esp. hepatitis and hepato-
                                                                                                Complementary; 5' untranslated region; UTR; hepatitis C virus; HCV; inhibition; replication; expression; detection; chronic hepatitis; acute hepatitis; hepatocarcinoma; ss.
                                                                                                                                                                                                                                                                                                                             /*tag=' b
/note= "Comprises phosphorothioate linkages"
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                                                         Oligo HCV-222, multiplex forming oligomer.
                                                                                                                                                                                                                      Location/Qualifiers
1 11
/ttag= a
7 notes 2 - OMe RNA"
12 . 16
/ttag= b
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hamlin HA,
Wolfe JL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (HOFF ) HOFFMANN LA ROCHE & CO AG F. (HYBR-) HYBRIDON INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 28; Page 22; 100pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 199 CCCTCCCGGGGACCTG 214
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                 (first entry)
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nes 13; Conservative
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, Walther DM,
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                                                                                                                                                                                                                             Key
modified_base
                                                                                                                                                                                                                                                                                                          modified_base
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Roberts PC,
                   16-OCT-1997
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                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       carcinoma
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule, that modulates human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule encoding HER2, K-Ras, H-Ras, N-Ras, human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-rheumatic activity. The nucleic acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung, prostate, bladder, or pancreatic cancer, and HIV infection, and AIDS. The sequences ABZ66530 - ABZ66585 represent substrate/target sequences for the human ribozymes of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences.
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                                                                                                                                                                                                                                                                                                                             Human, ribozyme; short interfering RNA, siRNA, HER2; K-Ras;
enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV;
anti-rheumatic; cancer; AIDS; ss.
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100.0%; Pred. No. 3.9e+02;
ive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                       Human H-Ras DNAzyme target #867.
                                                                                                                                                              ABZ62076 standard; RNA; 17 BP.
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06-JUN-2001; 2001US-0296249P.
10-SEP-2001; 2001US-0318471P.
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                 2 GGTTGCGGAGGGT 14
                                          GGTTGCGGAGGGT 4
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Matches 13, Conservative
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Roberts NA;

Kilkuskie RE,

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Gapa

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Indels

Query Match

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RESULT 413

AAT80369

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(KILK/)
                                                                                           (HAML/)
(ROBE/)
                                                                                                                 (WALT/)
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                                                      FRAN/
                                                                                   (ROBE/
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                                                                                                                                                                                                                                                                                                 The present invention relates to an oligonucleotide array comprising oligonucleotide tags fixed to a solid substrate. The oligonucleotide array is useful for genotyping a nucleic acid sample at one or more loci via single base extension (SBE) reactions. A pair of primers is used to amplify a polymorphic locus in a sample e.g. a single nucleotide polymorphism (SNP). The present sequence is one of the primers used in the method of the present invention to amplify a polymorphic sample. The amplified nucleic acid product is then used as a template in a SBE reaction with an extension primer. The SBE reaction products are used to
           Oligonucleotide array; genotyping; single base extension reaction; SBE; PCR primer; polymorphic locus; single nucleotide polymorphism; ss.
                                                                                                                                                                                                                                       Universal array of oligonucleotides tags attached to a solid substrate along with locus-specific tagged oligonucleotides useful in genotyping using single base extension reactions.
                                                                                                                                                                                         Б,
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                       Kaplan P, Lander ES, Lockhart
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hepatitis C virus; HCV; hepatocyte infection; non-A hepatitis; non-B hepatitis; acute hepatitis; chronic hepatitis; hepatocellular carcinoma; virucide; cytostatic; antisense therapy; gene therapy; ss; DNa-NRA hybrid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              .;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Inhibitory oligonucleotide specific for hepatitis C virus #159
                                                                                                                                                                                                                                                                                                                                                                                                                                        2.8%; Score 12.8; DB 1; Length 16; 87.5%; Pred. No. 3.8e+02; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 16 BP; 2 A; 2 C; 8 G; 4 T; 0 U; 0 Other;
                                                                                                                                                        (WHED ) WHITEHEAD INST BIOMEDICAL RES
                                                                                                                                                                                                                                                                                Example 7; Page 63; 70pp; English.
                                                                                                                                                                                       Huang X,
                                                                                                                                                                                                                                                                                                                                                                                                  form the oligonucleotide array
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP
                                                                                                     27-MAR-2000; 2000WO-US008069.
                                                                                                                         99US-0126473P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       97US-00887505.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24 AGGGGTGGTGGCCATT 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AGGGTTGGTGGCCAGT 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABS65953 standard; DNA; 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                   (AFFY-) AFFYMETRIX INC.
                                                                                                                                                                                      Hirschhorn JN,
                                                                                                                                                                                                                     WPI; 2000-656171/63
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                 Ryder T, Sklar P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US2002081577-A1
                                                             WO200058516-A2
                                          Unidentified.
                                                                                                                         26-MAR-1999;
23-JUN-1999;
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                                                                                 05-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15-NOV-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                             14;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABS65953;
                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                      Fan J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 415
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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The invention describes synthetic oligonucleotides complementary to a portion of the 5' untranslated region of hepatitis C virus. The oligonucleotides may be used in methods for controlling, preventing, and treating hepatitis C virus infection, in antisense technology and gene therapy, and of detecting the presence of hepatitis C virus in a sample. Hepatitis C virus (HCV) is an enveloped, positive sense, single-stranded RNA virus which infects hepatocytes. HCV is the major cause of non-A, non hepatocellular carcinoma. The invention describes methods and kits for inhibiting replication of HCV, inhibiting the expression of HCV nucleic acid and protein, and for treating HCV infections. This sequence represents a synthetic DNA-RNA hybrid oligonucleotide used for inhibiting HCV replication and expression of HCV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               useful for diagnosing and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eating disorder; polymorphism; dataset; allele; HGBASE identification; serotonin receptor 1D; delta-opioid receptor; dopamine receptor D2; anorexia nervosa; bulimia nervosa; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                                                                             Roberts PC;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic oligonucleotides complementary to a portion of untranslated region of hepatitis C virus (HCV), useful fo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               untranslated region of hepatitis C virus (HCV), usefu
treating HCV infections and hepatocellular carcinoma.
                                                                                                                                                                                                                                                                                                                                             Wolfe JL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 16 BP; 0 A; 7 C; 6 G; 2 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Serotonin receptor 1D probe SEQ ID No 61.
                                                                                                                                                                                                                                                                                                                                          Goodchild J,
Walther DM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 31; Page 7; 74pp; English.
95US-00471968.
96US-0021104P.
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20-JUL-2001; 2001US-0306440P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 2.8
Best Local Similarity 81.2
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                               Kilkuskie RL, Fraum
                                                                                  KILKUSKIE R L.
                                                                                                          FRANK B L.
GOODCHILD J.
WOLFE J L.
ROBERTS P C.
HAMLIN H A.
ROBERTS N A.
                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2002-537132/57.
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06-JUN-1995;
02-JUL-1996;
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Huntington's disease.
                                                                                                                                                                                                            invention.
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                                                                                                                                                                                                                                                                                                                                                        RESULT 418
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                                                                                                                                                                                 The invention relates to a novel isolated nucleic acid molecule comprising a variant gene associated with an eating disorder and selected from any of 119 polymorphisms with their corresponding genotyphism in the dataset, alleles and HGBASE identification, given in the specification. The novel nucleic acid molecule has polymorphisms in the serotonin receptor ID, delta-opioid receptor, or dopamine receptor D2, which is useful in diagnostic and prognostic assays for eating disorders, in particular ancreaxia nervosa and bulimia nervosa. This polymucleotide sequence represents a serotonin receptor ID probe of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New substantially pure ribonucleic acid (RNA) complex comprising a first strand and a second strand that hybridize to each other, useful for treating cancer, systemic lupus erythematosus, Alzheimer's disease or
                                                                                                         New nucleic acid molecule having polymorphisms in the serotonin receptor 1D, delta-opioid receptor, or dopamine receptor D2, useful in diagnostic and prognostic assays for eating disorders, such as anorexia and bulimia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   98; RNA complex; immunosuppressive; cytostatic; cancer;
systemic lupus erythematosus; Alzheimer's; Huntington's disease;
salivary gland carcinoma; melanoma; brain tumour; leukaemia; lymphoma;
                                                                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                                                                  Sequence 16 BP; 4 A; 10 C; 2 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mccallus DE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pyrimidine nucleotide flanking sequence 3.
                                                                                                                                                               Example 3; Page 60; 149pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                               ВЪ
13-NOV-2001; 2001US-0331285P.
19-DEC-2001; 2001US-0340843P.
19-DEC-2001; 2001US-0340844P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18-OCT-2002; 2002US-0419532P.
28-OCT-2002; 2002US-0421757P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-OCT-2003; 2003WO-US033466
                                                                                                                                                                                                                                                                                                                                                                   GGTTGCGGAGGGTGGG 17
                                                                                                                                                                                                                                                                                                                                                                               ADN14388 standard; DNA; 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                          (PRIC-) PRICE FOUND LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (NUCL-) NUCLEONICS INC.
                                                                Bergen AW, Yeager M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2004-348454/32.
                                                                                    WPI; 2003-268122/26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO2004035765-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      gene therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15-JUL-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                                                                                                                                                                                                   ~
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADN14388;
                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                            nervosa
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This invention relates to double stranded ribonucleic acid (RNA)

Structures and constructs. Specifically, it comprises first and second

Structures and constructs. Specifically, it comprises first and second

EXA strands that hybridise to each other under physiological conditions

to form a double-strand region, wherein the double-strand region contains

one or more mismatched regions that result in two or more double-stranded

segments. Furthermore, the mismatched regions may be cleaved by single-

strand ribonuclease enzymes. The present invention describes expression

vectors that encode daRNAs with an intron containing exemplary target

cetrois that encode daRNAs with an intron containing exemplary target

therapy, these RNA complexes exhibit immunosuppressive and cytostatic

cetivities and can be used to treat cancer, systemic lupus erythematosus,

Alzheimer's and Huntington's disease. The cancer is selected from,

melanoma, brain tumour, leukaemia and lymphoma. This oligonuclectide

sequence is a DNA flanking sequence used in an exemplification of the
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Antiviral phosphoro:thioate oligo:nucleotide(s) - active against e.g. herpes simplex virus 1, HIV and varicella zoster virus.
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/note= "phosphorothioate oligonucleotides"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 16 BP; 0 A; 10 C; 0 G; 6 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Mismatches
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Example 9; Page 119; 204pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             370 GAAGAGGAACGGAGCG 385
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94JP-00268603
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(KAKE ) KAKEN PHARM CO LTD.
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01-NOV-1994;
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RESULT 420
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                                                                                                                                                                                                                                                                                                                                       Antiviral; phosphorothioate; mRNA 4; mRNA 5; herpes simplex virus 1; HSV; viral infection; HIV; varicella zoster virus; VZV; therapy; ss.
            antiviral activity. These sequences, and the phosphorothioate oligonuclectides represented by AAT12418-T12434 (which are complementary to regions of the mRNA 4 or 5 of herpes simplex virus I (HSV)), are effective in the prevention and treatment of viral infection. The sequences are especially effective against infection by HSV, HIV or varicella zoster virus (VZV)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                antiviral activity. These sequences, and the phosphorothioate oligonuclectides represented by AAT12418-T12434 (which are complementary to regions of the mRNA 4 or 5 of herpes simplex virus 1 (HSV), are effective in the prevention and treatment of viral infection. The sequences are especially effective against infection by HSV, HIV or
                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                against e.g
AAT12435-T12454 represent phosphorothioate oligonucleotides with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAT12435-T12454 represent phosphorothioate oligonucleotides with
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                                                                                                                2.8%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 4e+02; tive 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                   /note= "phosphorothioate oligonucleotides"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Antiviral phosphoro:thioate oligo:nucleotide(s) - active herpes simplex virus 1, HIV and varicella zoster virus.
                                                                                           Sequence 17 BP; 0 A; 2 C; 15 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 17 BP; 0 A; 3 C; 14 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                  Antiviral phosphorothioate oligonucleotide #26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Iwatani W,
                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Shimada J, Mizushima Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 6; Page 150; 163pp; Japanese.
                                                                                                                                                                                                                                                ВР
                                                                                                                                                               333 GGGGGGGAGGCGAGG 348
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (LTTL-) LTT INST CO LTD. (KAKE ) KAKEN PHARM CO LTD.
                                                                                                                                                                                                                                               AAT12443 standard; DNA; 17
                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                              Query Match
Best Local Similarity 87.5
Matches 14; Conservative
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Best Local Similarity
Matches 14; Conserv
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                                                                                                      AAT12443;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Shoji Y,
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The present invention describes an enzymatic nucleic acid molecule (I) with RNA cleaving activity, which modulates the expression of a plant gene. Also described is a gene comprising a cDNA sequence encoding maize Delta-9 desaturase. (I) can be used to modulate expression of a gene, preferably Delta-9 desaturase or a granule bound starch synthase (GBSS) gene, in a plant (preferably a maize or canola plant). (I) can be used to modulate caffeins synthesis in a coffee plant, nicotine production in a tobacco plant, fruit ripening processes in an apple, tomato, pear, plum or peach plant, flower pigmentation in a rose, petunia, chrysanthemum or marigold plant or lignin production in a tobacco, aspen, poplar or pine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Maize; corn; Zea mays; delta-9 desaturase; GBSS; target; substrate; granule bound starch synthase; hammerhead ribozyme; hairpin ribozyme; modulation; gene expression; transgenic plant; cleavage; canola plant; caffeine synthesis; coffee plant; nicotine production; tobacco; fruit ripening; flower pigmentation; lignin production; se.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Skokut TA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        expression - preferably modulates granule bound starch synthase in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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Merlo DJ;
                                                                                                                                                                                                                                                                                                                                                                                                              Delta-9 desaturase hamerhead ribozyme target SEQ ID NO:828.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2.8%; Score 12.8; DB 1; Length 17; 2.5%; Pred. No. 4e+02; ve 4; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ribozyme which modulates plant gene expression of DELTA-9 desaturase or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 38; Page 86; 155pp; English.
332 CGGGGGGGGGGGGGG 347
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                                                        2 cededecedecede 17
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nes 10; Conservative
                                                                                                                                                                                                                      AAX62953 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Edington BE,
Folkerts O,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1997-202224/18.
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Young SA,
                                                                                                                                                                                                                                                                                 AAX62953;
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WPI; 1999-047888/05
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        30-DEC-1998
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                                                                                Blasczyk R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAX00304;
                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 423
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                                                                                                                                                                                                                                                                 Detection of mutation(s) in the BRCAl gene - by hybridisation with an allele-specific oligo:nucleotide or by amplification, useful particularly for breast or ovarian cancers.
                                                                                                                                                                                                                                                                                                                      AAV20567-V20574 are probes used in a method to detect mutations in the ... human BRCA1 gene. Such mutations are used for classifying a tumour for diagnostic and prognostic purposes or detecting a predisposition of higher susceptibility to breast and ovarian cancer in an individual. The methods can be used for reducing the high incidence and mortality women at high risk. These women, once identified, can be targeted for more aggressive prevention programmes
                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human Leukocyte Antigen; HLA; HLA-DRB consensus sequence; intron 1;
HLA class II group type; histocompatibility analysis;
compatibility analysis; PCR primer; 88.
                                                   Breast cancer; ovarian cancer; mutation; classification; detection; tumour; diagnostic; prognostic; probe; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                           2.8%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 4e+02; ive 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                Olson SJ,
                                                                                                                                                                                                                                                                                                        Example 4; Page 40; 62pp; English.
                                                                                                                                                                                                                                White MB,
                                                                                                                                                               96US-0023184P.
96US-0023187P.
96US-002323P.
96US-0022421P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 366 GCAGGAAGAGGAACGG 381
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                  02-JUL-1998 (first entry)
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Best Local Similarity 87.5
Matches 14; Conservative
                                                                                                                                                                                                                                Murphy PD, Allen AC,
                                   Human BRCAl probe #4.
                                                                                                                                                                                                              (ONCO-) ONCORMED INC.
                                                                                                                                                                                                                                                WPI; 1998-159166/14.
                                                                                          Homo sapiens.
                                                                                                                                                                                  05-AUG-1996;
06-AUG-1996;
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                                                                                                           WO9805677-A1
                                                                                                                                               04-AUG-1997;
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05-AUG-1996;
                                                                                                                             12-FEB-1998
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                                                                                Synthetic
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AAV20570;
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Leukocyte Antigen (HLA)-DRB sequences. The primers are used in the methods of the invention. The specification describes a method for determining the HLA class II group type of a subject. The method for determining the HLA class II group type of a subject. The method comprises amplifying a target DNA sample from a subject using a particular HLA group-specific primer pair and determining whether a nucleic acid product is produced, therefore identifying the group type. Methods for determining the HLA class allele type of a subject are also described, where a specific HLA group-specific exon region primer pair is used. The methods are useful for determining the HLA class II type of a getermining the group precific alleles present and determining the group specificatly of alleles. The methods are diagnostically useful for histocompatibility analysis
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                                                                                                                                                                                                                                                                                                                                                                   Determining the Human Leukocyte Antigen Class II type Histocompatibility antigens - by using new intron-specific oligonucleotide primers for sequence specific primer PCR and sequencing.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2.8%; Score 12.8; DB 17.5%; Pred. No. 4e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 13; Fig 3A; 36pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         232 AGCCCCCGAACCCCGC 247
97EP-00110438
                                                                         97EP-00110438
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Best Local Similarity 87.5%;
Matches 14; Conservative
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class II (HLA Class II) group type of a subject the method comprises:

(i) amplifying a target DNA sample from a subject using a particular HLA group-specific primer pair (sequence specific primer PCR - SSP-PCR); and group-specific primer pair (sequence specific primer PCR - SSP-PCR); and claimed by the group type. AAX00305 to AAX00305 tepresent specifically claimed oligonucleotide primer for use in the above method. These oligonucleotides are useful for determining the HLA Class II type of a patient sample, by identifying the specific alleles present and etermining the group specific alleles. Steps (i) and (ii) in the method are diagnostically useful for histocompatibility analysis to see if donor and recipient groups match. The new sequences are useful for providing an insight into the genetic relationship between different angency of HLA Class II genes. The high resolution, nucleic acid based primer be indight the intron-specific primers is more efficient than prior art methods using exon based primers, as few exon sequences offer conserved primer binding sites, resulting in a limited number of primer pairs and the primer specificity for alleles, as allelic variations exist between the primer sites. The SSP-PCR method allows separation of haplotypes in the primer sites. The SSP-PCR method allows separation of cis-trans linkages of partences are useful for any and the primer sites. The sylventy cannot be achieved with other primers.
                                    Determining the Human Leukocyte Antigen Class II type Histocompatibility antigens - by using new intron-specific oligonucleotide primers for sequence specific primer PCR and sequencing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Antisense oligonucleotide, multiple target; antisense treatment; impaired respiration; inflammation; lung disease; pulmonary vasconstriction; inflammation; allergic rhinitis; acute asthma; allergy; asthma; impeded respiration; respiratory distress syndrome; pain; cystic fibrosis; pulmonary hypertension; pulmonary vasconstriction; emphysema; chronic obstructive pulmonary disease; leukemia; lymphoma; carcinoma; colon cancer; breast cancer; lung cancer; pancreatic cancer; hepatocellular carcinoma; kidney cancer; melanoma; hepatic metastasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2.8%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 4e+02; tive 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C/EBP-beta antisense oligonucleotide fragment.
                                                                                                                               Claim 4; Fig 3; 36pp; English
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98US-00093972.
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WPI; 1999-083585/08
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09-JUN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      protocols
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAX55127;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 424
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The specification describes antisense oligonucleotides (AAX52869-X55271)

directed against at least 2 mRNAs selected from target genes, coding and
non-coding regions of RNAs corresponding to target genes, coding and
codons, genomic flanking regions, intron-exon borders, the 5'-end, the 3'
codons, genomic flanking regions, intron-exon borders, the 5'-end, the 3'
codons, genomic flanking regions, intron-exon borders, the 6'-end, the 3'
codons, mixtures. The antisense oligonucleotides may be derived
conditions or mixtures. The antisense oligonucleotides may be derived
from sequences AAX5512-74. These multiple target oligonucleotides
conditions. Typical diseases and conditions are those
associated with impaired respiration and inflammation, including lung
diseases, pulmonary vasoconstriction, inflammation, allergic rhinitis,
acute asthma, allergies, asthma, impeded respiration, respiratory
diseases (COPD), and cancers such as leukemias, lymphomas, carcinomas e.g.
colon cancer, breast cancer, lung cancer, pancreatic cancer,
colon cancer, breast cancer, lung cancer, pancreatic metastassized
colon cancer, breast cancer, lung cancer, pancreatic cancer,
chepatocellular carcinoma, kidney cancer, metastassized
colon cancer, breast cancer, mine and prostate cancer
colon cancer, breast cancer, lung cancer, pancrealic cancer,
colon cancer, breast cancer, lung cancer, pancrealic cancer,
colon cancer, breast cancer, metastasized
colon cancer, breast cancer, metastasized ö phosphorothioate; impaired respiration; inflammation; allergy; allergy; allergic disease; bronchoconstriction; inhibitor; antiinflammatory; antiallergic; antiasthmatic; cytostatic; analgesic; impaired airway; lung disease; ischaemic condition; pulmonary vasoconstriction; asthma; respiratory distress syndrome; pain; cystic fibrosis; emphysema; pulmonary hypertension; chronic obstructive pulmonary disease; COPD; Gaps New antisense oligonucleotides used in treatment of, e.g. pulmonary Human; adenosine receptor; low adenosine antisense oligonucleotide; ; 0 Human adenosine receptor related polynucleotide SEQ ID NO:2263 2.8%; Score 12.8; DB 1; Length 17; Local Similarity 87.5%; Pred. No. 4e+02; es 14; Conservative 0; Mismarchor cancer; leukaemia; lymphoma; carcinoma; metastasis; ss. Sequence 17 BP; 0 A; 12 C; 5 G; 0 T; 0 U; 0 Other; Disclosure; Page 71; 120pp; English. 272 BP 99WO-US017712. 98US-0095212P. (UYEC-) UNIV EAST CAROLINA. AAA34574 standard; DNA; 17 28-JUL-2000 (first entry) 257 GCGGTCGGCCCGGGGC 16 gcagacacacacagac WPI; 1999-229400/19 vasoconstriction WO200009525-A2. sapiens. 03-AUG-1999; 03-AUG-1998; 24-FEB-2000 AAA34574; Query Match Nyce JW; Best Loca Matches RESULT 425 Homo AAA34574/ ò

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Gaps . 0 Low adenosine (A) content antisense oligonucleotides which do not trigger adenosine receptors during metabolism, useful e.g. for treating cancers

Claim 14; Page 265; 1592pp; English

and respiratory obstructions

WPI; 2000-679539/66.

24-MAR-2000; 2000WO-US008020

WO200062736-A2

26-OCT-2000

99US-0127958P.

06-APR-1999;

(UYEC-) UNIV EAST CAROLINA.

(NYCE/) NYCE J W.

Nyce JW;

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                                                                                                                                                                                                                                                                                                                                                             The present invention describes a new composition comprising an antisense oligonucleotide (ON) with low adenosine (up to 15%), which targets concletic acids involved in bronchoconscriction, allergies, and/or inflammation. The ON can have antinflammatory, antiallergic, antiasthmatic, cytostatic and analgesic activities. The compositions are useful for the treatment of diseases associated with inflammation, impaired airways, including lung disease associated with inflammation, impaired airways, including lung disease and diseases whose secondary effects afflict the lungs of a subject. They can be used for treating c. g. ischaemic conditions, pulmonary vasoconstriction, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, pulmonary hypertension, emphysema, chronic obstructive carcinomas, and cancers which may metastasis to the lungs, including breast and prostate cancer. The reduction of the adenosine content of the carcinomas, and effects. The A-containing ONs break down with the release of deoxyadenosine which activates adenosine receptors causing conchoconstriction and inflammation. AAA3313 to AAA3312 represent the nucleotide sequences given in the sequence listing from the present invention, which correspond to SEQ ID No:11 to 1885 from the present invention. N.B. Sequences given in the disclosure of the present invention on the disclosure of the present invention of onct match is the interest given in the disclosure of the present invention do not match is the interest given in the sequences given in the sequences given in the sequences given in the sequences given in the esquences given in the disclosure of the present invention do not match the interest and prostates and prostates given in the disclosure of the present invention do not match the sequences given in the disclosure of the greatest sequences given in the disclosure of the greatest sequences.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                    New antisense oligonucleotides useful for treating e.g. pulmonary vasoconstruction, inflammation, allergies, asthma, hypertension, bronchitis, emphysema, respiratory distress syndrome, ischemia or
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87.5%; Pred. No. 4e+02;
tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 17 BP; 0 A; 12 C; 5 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human C/EBP polynucleotide fragment #2263.
                                                                                                                                                                                                                                                                                                                   Disclosure; Page 548; 1343pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAF20696 standard; DNA; 17 BP.
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(UYEC-) UNIV EAST CAROLINA
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                                                                                                             WPI; 2000-205971/18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
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                                                         Nyce JW
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The present invention describes low authoring why content and the antisense oligonucleotides and compositions (I) comprising them. In the antisense oligonucleotides the A is replaced by a 'Universal' or alternative base.

(I) can have respiratory, bronchodilator, antihiflammatory, analgeat, immunosuppressive, antiaethmatic, hypotensive and cytostatic activities. The antisense oligonucleotides and (I) can be used to down-regulate the expression and or activity of target polypeptides associated with activating peptide factors and malignancies, auch as stimulating and activating peptide factors and transmitters, transcription factors, immunoglobulins and antibodies, antibody receptors, cytokines and chemokines, endogenously produced specific and non-apecific anzymes, chemokines, addesion molecules and their receptors, central nervous and non-nervous system control nervous system (CNS) and peripheral nervous and non-nervous system peptide transmitters, defensins, growth factors, vasocative peptides and receptors, binding proteins and malignancy associated proteins. The receptors, binding proteins and malignancy associated disorders including respiratory obstruction (especially pulmonary obstruction and/or bronchoconstriction) and/or lung inflammation, allergy(ies) and/or condition and formulance associated with a disease or condition and formulance associated with a disease or condition and formulance associated with a disease or conditions.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     condition selected from pulmonary vasoconstriction, inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome (RDS), pain, cystic fibrosis (CF), allergic rhinitis (AR), pulmonary hypertension, emphysema, chronic obstructive pulmonary disease (COPD), pulmonary transplantation rejection, pulmonary infections, bronchitis, and/or cancer. AAF18434 to AAF21543 represent human polynucleotide fragments and antisense oligonucleotides used in the exemplification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2.8%; Score 12.8; DB 1; Length 17;
87.5%; Pred. No. 4e+02;
tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 17 BP; 0 A; 12 C; 5 G; 0 T; 0 U; 0 Other;
present invention describes low adenosine (A)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hammerhead ribozyme substrate #2955.
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Best Local Similarity 87.5
Matches 14; Conservative
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The invention relates to the construction of a vector which comprises a multiple number of Shear Stress Response Elements (SSRE) from various can gene promoter sequences and one or more genes, antisense molecules, cribozymes, double stranded RNA, or a nucleic acid which encodes a cribozymes, double stranded RNA, or a nucleic acid which encodes a cribozymes, double stranded RNA, or a nucleic acid which encodes a control of the protein which inhibits the synthesis of, or activity of the protein or peptide. This sequence represents the SSRE sequence from the promoter of the platelet-derived growth factor A (PDGF-CC A). The vector is useful for stimulating or inhibiting vascular permeability or seen of interest is custing the formation, maturation or regression of blood vessels, or inhibiting genes or proteins involved in a diseases, down regulating angiogenesis and for treating vasculogenic angiogenesis and for treating vasculogenic angiogenesis. These disorders include cardiovascular disorder, neoplastic disorders, is chaemia, atheresclerosis, hypertension, diabetes, hypercholesterolaemia
                                                                                                            Expression vector comprising multiple shear stress response elements, useful for modulating endothelial cell proliferation, stimulating or down-regulating angiogenesis and treating vasculogenic/angiogenic disorders.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             checkpoint kinase-1; Chk1; antisense; ribozyme; gene therapy;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ouery Match 2.8%; Score 12.8; DB 1; Length 17; Best Local Similarity 87.5%; Pred. No. 4e+02; Matches 14; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 17 BP; 0 A; 2 C; 15 G; 0 T; 0 U; 0 Other;
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                                                                                                                                                                                                            Example 1; Page 45; 61pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    333 GGGGCGAGGGCGAGG 348
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                                                                     WPI; 2000-452382/39.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        and wound healing
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                        Resnick N;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to enzymatic and antisense nucleic acid molecules that act as inhibitors of the expression of repressor genes encoding the TR2 Orphan receptor, EAR3/COUP-TF-1, the GATA transcription factor gene, IRF-2 and/or the CAATT Displacement Protein (CDP). Inhibition of the repressors removes prevents inhibition (and consequently increases expression of) genes involved in the production of erythropoietin, granulocyte colony stimulating factor protein and
                                                                                                                                                                                                                                                                                                                                                                                                                         Enzymatic and antisense nucleic acid inhibition of repressor genes, useful for producing e.g. granulocyte colony stimulating factor protein, interferon alpha and erythropoietin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cytostatic, cardiant, vasotropic, vulnerary, antidiabetic, hypotensive, antiatherosclerotic, antilipemic, gene therapy, vector, SSRE, promoter, Shear Stress Response Element, antisense, riboxyme, repressor antibody, platelet derived growth factor A, PDGF-A, angiogenesis, ischaemia, cardiovascular disorder, neoplastic disorder, atherosclerosis, ss;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          hypertension; diabetes; hypercholesterolaemia; wound healing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2.8%; Score 12.8; DB 1; Length 17; 37.5%; Pred. No. 4e+02; rative 8; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Shear Stress Response Element from PGDF-A gene.
                                                                                                                                                                                                                                                                                                                                     Mcswiggen J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 42; Page 123; 164pp; English.
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                                                                                                                                                                                         11-APR-2000; 2000WO-US009721.
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                                                                                                                                                                                                                                                                                 (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
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interferon alpha; ss
                                                                                                                                                                                                                                                                                                                                                                             WPI; 2000-647423/62.
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Les 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         interferon alpha
                                                                                           WO200061729-A2
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                                                 Homo sapiens.
                                                                                                                                                                                                                                       12-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      23-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           24-DEC-1998;
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                                                                                                                                          19-OCT-2000
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Gaps

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Claim 4; Page 52; 115pp; English

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BP

(first entry)

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ABK00059 standard; RNA; 17
                                                    12-MAR-2002
                                ABK00059;
ABK00059/
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention provides nucleic acid molecules capable of downregulating the expression of the human checkpoint kinase-1 (Chkl) gene. These may be antisense or ribozyme sequences, and are useful in the treatment of diseases associated with conditions affected by Chkl levels, including cancer. The present sequence is an oligonucleotide described in the exemplification of the invention
         downregulating the expression of the human checkpoint kinase-1 (Chk1) gene. These may be antisense or ribozyme sequences, and are useful in the treatment of diseases associated with conditions affected by Chk1 levels, including cancer. The present sequence is an oligonucleotide described in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel nucleic acid molecule e.g., ribozymes or antisense nucleic acid molecules, which downregulates expression of a checkpoint kinase-1 geruseful for treating colorectal, lung, breast or prostate cancers.
                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                   Human; checkpoint kinase-1; Chk1; antisense; ribozyme; gene therapy;
The present invention provides nucleic acid molecules capable of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Booher RN, Holman PS;
                                                                                               ch 2.8%; Score 12.8; DB 1; Length 17; I Similarity 87.5%; Pred. No. 4e+02; 14; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 2.8%; Score 12.8; DB 1; Length 17; Best Local Similarity 87.5%; Pred. No. 4e+02;
                                                                           Sequence 17 BP; 5 A; 2 C; 7 G; 0 T; 3 U; 0 Other;
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                                                                                                                                                                                                                                                                                            Human Chkl ribozyme substrate SEQ ID NO: 959.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mcswiggen J,
                                                      the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 4; Page 77; 115pp; English
                                                                                                                                                                                                                         AAH95534 standard; RNA; 17 BP
                                                                                                                                              272 CTTCTCCGGAGGCACC 287
                                                                                                                                                                                                                                                                                                                                                                                                                      02-FEB-2001; 2001WO-US003504
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(FATT/) FATTAEY A R.
                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                               RNA cleavage; cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Jarvis T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-496922/54.
                                                                                                                                                                                                                                                                                                                                                                         WO200157206-A2.
                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
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Best Local 3
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The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down cregulates expression of a neutite growth inhibitor gene (NGGO). The nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a DNAzyme) an Inozyme (an endolytic nucleic acid cleaving an RNA molecule possessing an NCH motif), a G-cleaver (cleaving RNA with a NYN motif) properties or a maberzyme (cleaving RNA with an NGN triplet), a zinzyme (cleaving RNA with a YGY motif). The CD20-targeting nucleic acid is used to cleave RNA cross of CD20 in the presence of a divalent cation that is preferably Mg^2+.

Furthermore, it may be contacted with a cell to reduce CD20 activity of the cell and treat a patient having a condition associated with the level of CD20. The treatment may further comprise the use of one or more characters. In particular, the CD20 targeting nucleic acid may be used to treat lymphoma, leukaemia, B-cell lymphoma, low-grade or follicular non-thody inmunnocytoma (NHL), immunocytoma (NHL), immunocytoma (NHL), immunocytoma (MCL), imm
                                                                                              Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; nootropic; neuroprotective; antiparkinsonlan; muscular; CD20; neurite growth inhibitor gene; NOGO; hammerhead ribozyme; DNAzyme; incygme; G-cleaver; amberzyme; inzyme; Juymphoma; leukaemia; B-cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphocytic leukaemia; human immunodeficiency virus; HIV associated NHL; nantle-cell lymphoma; MCI; immunocytoma; IMC; immune thrombocytopaenia; stroke; dementia; inflammatory archropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis; chemotherapy-induced neuropathy; amyotrophic lateral sclerosis; ALS; Parkinson's disease; ataxia; Huntington's disease; creutzfeldt-Jakob disease; muscular dystrophy; neurodegenerative disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chowrira BM;
Human NOGO Hammerhead Ribozyme #59.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 88; Page 66; 200pp; English.
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28-FEB-2000; 2000US-0185516P.
06-MAR-2000; 2000US-0187128P.
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Synthetic.
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2; Indels

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14; Conservative

Matches

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17 criciccaragecace

Claim 88; Page 66; 200pp; English.

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presence of a divalent cation that is preferably Mg^2^+. Furthermore, the nucleic acid may be contacted with a cell to reduce NOGO activity of the cell and treat a patient having a condition associated with the level of NOGO. The treatment may further comprise the use of one or more therapies. In particular, the NOGO-targetting nucleic acid may be used to treat central nervous system (CNS) injury and cerebrovascular accident (CVA, stroke), Alzheimer's disease, dementia, multiple sclerosis (MS), chemotherapy-induced neuropathy, amyotrophic lateral sclerosis (MS), darkinson's disease, ataxia, Huntington's disease, Creutzfeldt-Jakob disease, muscular dystrophy, and/or other neurodegenerative disease sequence is a hammerhead ribozyme of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 17 BP; 1 A; 6 C; 3 G; 0 T; 7 U; 0 Other;
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28-FEB-2000; 2000US-0185516P.
06-MAR-2000; 2000US-0187128P.
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BLATT L.
MCSWIGGEN J.
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(MCSW/) 1
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The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (NGO). The regulates expression of a neurite growth inhibitor gene (NGO). The concleic acids may be enzymatic nucleic acid cleaving a an RNA molecule possessing an NCH motifi), a G-cleaver (cleaving RNA with a NYM motif), a G-cleaver (cleaving RNA with a NYM motif) proposessing an NCH motifi). The CD20-targetting nucleic acid is used to cleave RNA of CD20 in the presence of a divalent cation that is preferably MG<sup>2</sup>+.

CC with a YGY motif). The CD20-targetting nucleic acid is used to cleave RNA of CD20 in the presence of a divalent cation that is preferably MG<sup>2</sup>+.

CC FURTHERMORE, it may be contacted with a call to reduce CD20 activity of the CC treat lymphoma, leukaemia, B-cell lymphoma, low-grade or follicular non-clear lymphoma, leukaemia, B-cell lymphoma, low-grade or follicular non-cleaker lymphoma (MCL), immunocytoma (IMC), small B-cell lymphocytic lymphoma (MCL), immunocytoma (IMC), small B-cell lymphocytic lymphoma, leukaemia, HIV (human immunodeficiency virus) associated MHL, mantle-cell lymphoma (MCL), immunocytoma (IMC), small B-cell lymphocytic lymphoma (MCL), immunocytoma (IMC), small secolated with the level of tragetting nucleic acid may be contacted with a cell to reduce NGOs activity of the collicia call may be used to contacted with a cell to reduce NGOs activity of the collicia acid may be used to create central nervous system (CNS) injury and creates NGOs activity of second collicia call may be used to create central nervous system (CNS) injury and creates of one or more collinear central nervous system (CNS) injury and creates of creates which respond to the meurodegenerative creates which respond to the neurodegenerative 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; nootropic; neuroprotective; antiparkinsonian; muscular; D20; neurite growth inhibitor gene; NOGO; hammerhead ribozyme; DNAzyme; inozyme; g-cleaver; amberzyme; zinzyme; lymphoma; leukaemia; B-cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphocytic leukaemia; MCL; immunocytoma; non-Hodgkin's lymphoma; NHL; lymphocytic leukaemia; MCL; immunocytoma; IMC; immune thrombocytopaenia; stroke; dementia; inflammatory arthropathy; central nervous system injury; centebrotypascular accident; CVA; Alzheimer's disease; multiple sclerosis; chemotherapy-induced neuropathy; amyotrophic lateral sclerosis; ALS; parkinson's disease; ataxia; Huntington's disease; Creutzfeldt-Jakob disease; muscular dystrophy; neurodegenerative disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        268
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABK01810 standard; RNA; 17 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12-MAR-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  253 GGCCGCGGTCGGCCCG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16 GGCCGCGGACAGCCCG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human NOGO Zinzyme #132.
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WO200159103-A2

Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia, and central nervous system injury.

Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; nootropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NOG0; hammerhead ribozyme; DNAzyme; inozyme; dever; amberzyme; inozyme; lincyme; lincyme; lincyme; lincyme; lincyme; leukaemia; B-cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphom; leukaemia; B-cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphocytic leukaemia; MCL; immunocytoma; IMC; immune thrombocytopaenia; stroke; dementia; inflammatory arthropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis; chemotherapy-induced neuropathy; amyotrophic lateral sclerosis; ALS; Parkinson's disease; ataxia; Huntington's disease; cerebrovascular dystrophy; neurodegenerative disease.

Human NOGO Hammerhead Ribozyme #58.

12-MAR-2002 (first entry)

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The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down capuates expression of a neurite growth inhibitor gene (NGOD). The nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a DNAzyme) an Inozyme (an endolytic nucleic acid cleaving a an RNA molecule DNAzyme) an Inozyme (an endolytic nucleic acid cleaving an RNA molecule an amberzyme (Cleaving RNA with an NGW niple!), a zinzyme (cleaving RNA with an NGW triple!), a zinzyme (cleaving RNA with a cell low-grade or follicular non-leukaemia, HV (human immunodeficiency virus) associated NHL, immunocytoma (IMC), smail B-cell lymphomy (LI), immunocytoma (IMC), smail B-cell lymphocytic lymphomy (CC tragetion asystem (IMC), smail B-cell lymphocytic lymphomy (I
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          muscular dystrophy, and/or other neurodegenerative disease hich respond to the modulation of NOGO expression. The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 17 BP; 0 A; 7 C; 8 G; 0 T; 2 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          states which respond to the modulation of NOGO esquence is a zinzyme molecule of the invention
                                                                                                                                                                                                                                                                                                                                                     Chowrira BM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 88; Page 98; 200pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  central nervous system injury.
                                                                         09-FEB-2001; 2001WO-US004273.
                                                                                                                        2000US-0181797P
                                                                                                                                                28-FEB-2000; 2000US-0185516P.
06-MAR-2000; 2000US-0187128P.
                                                                                                                                                                                                                       RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                     Blatt L, Mcswiggen J,
                                                                                                                                                                                                                                                                      MCSWIGGEN J.
CHOWRIRA B M.
                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-607195/69.
                                                                                                                                                                                                                                                      BLATT L.
                                                                                                                        11-FEB-2000;
                    16-AUG-2001
                                                                                                                                                                                                                                                 (BLAT/)
(MCSW/)
(CHOW/)
                                                                                                                                                                                                                         (RIBO-)
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28-FEB-2000; 2000US-0185516P. 06-MAR-2000; 2000US-0187128P. 11-FEB-2000; 2000US-0181797P. 09-FEB-2001; 2001WO-US004273

WO200159103-A2.

16-AUG-2001

sapiens.

Ношо

Synthetic.

(RIBO-) RIBOZYME PHARM INC.

BLATT L.

(MCSW/) BLAT/)

Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia, and

Claim 88; Page 66; 200pp; English. central nervous system injury.

Chowrira BM;

Blatt L, Mcswiggen J, CHOWRIRA B M MCSWIGGEN J.

WPI; 2001-607195/69.

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                                   Gaps
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Query Match 2.8%; Score 12.8; DB 1; Length 17; Best Local Similarity 87.5%; Pred. No. 4e+02;
                                 2; Indels
                                 0; Mismatches
                                                                 254 GCCGCGGTCGGCCCGG 269
                                 14; Conservative
                                 Matches
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The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down cregulates expression of a neutite growth inhibitor gene (NDGO). The nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a DNAzyme) an Inozyme (an enadolytic nucleic acids (e.g. a ribozyme or a maberzyme). The CD20 tergetting nucleic acid is used to cleave RNA with a NYN mouth? properties and north mouth? The CD20 tergetting nucleic acid is used to cleave RNA with a YGY motif). The CD20 tergetting nucleic acid is used to cleave RNA vith a YGY motif). The CD20 tergetting nucleic acid is used to cleave RNA clear the reat a patient having a condition associated with the level of CD20. The treatment may further comprise the use of one or more therapies. In particular, the CD20 tergetting nucleic acid may be used to treat lymphoma, leukaemia, B-cell lymphoma, low-grade or follicular non-Hodgkin's lymphoma (NHL), bulky low-grade or follicular NHL, lymphocytic leukaemia, HV (human immunodeficiancy virus) associated NHL, mantle-cell lymphoma (MCL), immunocytoma (IMC), small B-cell lymphocytic lymphoma, immunodeficiancy virus) associated NHC, mantle-cell lymphoma (MCL), immunocytoma (IMC), small B-cell lymphocytic lymphoma, immunodeficiancy virus) associated NHC, mantle-cell lymphoma (MCL), immunocytoma (IMC), small B-cell lymphocytic lymphoma, immunodeficiancy virus) associated NHC, lymphoma, immunodeficiancy virus) associated vith the level of nucleic acid may be contacted with a cell to reduce Of one or more therapies. In particular, immunocytoma (CNS) injury and cerebrone of a l
17 GCCGCGGACAGCCCGG
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(CVA, stroke), Alzheimer's disease, dementia, multiple sclerosis (MS),

ABK00058 standard; RNA; 17 BP.

RESULT 434

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ABK00058

Human; UDP-GalNAc:polypeptide N-acetylgalactosaminyltransferase 10; pp-GaNTase 10; EC 2.4.1.41; chromosome 7q11.2; gene therapy; scanning;

Homo sapiens. Synthetic. EP1243660-A2.

Human pp-GaNTase 10 scanning 17-mer SEQ ID NO:738

(first entry)

11-DEC-2002

ABV85745;

BP.

ABV85745 standard; DNA; 17

RESULT 43(ABV85745/c

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           Chemotherapy-induced neuropathy, amyorrophic lateral sclerosis (ALS), Parkinson's disease, ataxia, Huntington's disease, returifieldt Jakob disease, muscular dystrophy, and/or other neurodegenerative disease states which respond to the modulation of NOGO expression. The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Domain
                                                                                                                                                                                                                                                                                                                                                                            Gaps
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co-stimulatory adaptor protein; tissue rejection; graft rejection;
leukaemia; cytostatic; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ellis JH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New nucleic acid(s) for regulating the Grb2-related with Insert D (GRID) gene comprises using antisense and enzymatic nucleic acid molecules such as hammerhead ribozymes.
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2.8%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 4e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human GRID NCH ribozyme substrate oligonucleotide #152.
chemotherapy-induced neuropathy, amyotrophic lateral
                                                                                                                                          sequence is a hammerhead ribozyme of the invention
                                                                                                                                                                                                                  Sequence 17 BP; 0 A; 6 C; 3 G; 0 T; 8 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                    363 GCCGCAGGAAGAGAA 378
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABL46698 standard; RNA; 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABL46698;
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30-JAN-2001, 2001WO-US000665. 30-JAN-2001, 2001WO-US000666. 30-JAN-2001, 2001WO-US000667. 30-JAN-2001, 2001WO-US000669. 30-JAN-2001, 2001WO-US000669. 31-JAN-2001, 2001WO-US000670. 23-WAY-2001, 2001WS-0315984F.

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Gu Y, Nguyen

Zhang J,

WPI; 2002-724954/79.

(AEOM-) AEOMICA INC.

25-JAN-2002; 2002EP-00001161

25-SEP-2002

2001WO-US000663 2001WO-US000664

30-JAN-2001;

30-JAN-2001;

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Nucleic acid encoding human UDP-GalNAc:polypeptide N-cetylgalactosaminyltransferase 10 protein is useful to diagnose, prevent and treat disorders associated with reduced or over expression of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention describes an isolated nucleic acid (I) encoding a human UDP-GalNAc:polypeptide N-acetylgalactosaminyltransferase 10 (pp-GaNTaee 10, EC 2.4.1.41) protein. Human pp-GaNTaee 10 is located to chromosome 7411.2. (I) can be used in gene therapy. Molecules of the present invention can be used in therapy, particularly to prevent or treat a disorder associated with decreased expression or activity of pp-GaNTaes. The sequences given in ABV85011 to ABV86689 and ABPS530 to The sequence data for this patent is not represented in the printed specification of the present invention. N.B. specification but is based on sequence information supplied by the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 17 BP; 3 A; 6 C; 6 G; 2 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       encoded protein.
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RESULT 437 ABV85746/c

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Gaps

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2.8%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 4e+02; tive 0; Mismatches 2; Indels

200 CCTCCCGGGACCTGC 215

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14; Conservative

Matches

Local Similarity

Query Match

16 ccrcccrcccicc 1

(first entry)

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Chromosomal genomic alteration; genome altering oligonucleotide; PCR; 88; o-methyl modification; LNA modification; phosphorothioate linkage; DNA repair; DNA alteration; environmental tolerance; hygromychnib phosomychnib abiotic stress tolerance; improved nutritional value; hygromychn; primer; amino acid over production; herbicide resistance; glyphosate resistance; porphyric herbicide resistance; sulphonylurea herbicide resistance; modified oil production; modified starch production; waxy starch; modified floral morphology; male-sterile plant; albino mutant; modified fatty acid content; reduced plant; albino mutant; increased stearate production; albino plant; increased stearate production; albino plant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New oligonucleotides with modified nuclease-resistant termini, useful for creating plants with desired phenotypes, e.g. stress tolerance, improved nutritional value, herbicide or disease resistance, or modified oil
                                                         Male-sterile plant producing genome altering oligonucleotide #291
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-JUN-2000; 2000US-0208538P.
30-OCT-2000; 2000US-0244989P.
27-MAR-2001; 2001US-00818875.
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                                                                                                                                                                                                                                                                                                                         photosynthetic process.
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                09-APR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     06-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
Nucleic acid encoding human UDP-GalNAc:polypeptide N-cetylgalactosaminyltransferase 10 protein is useful to diagnose, prevent and treat disorders associated with reduced or over expression of the encoded protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention describes an isolated nucleic acid (I) encoding a human UDP-GalNAc:polypeptide N-acetylgalactosaminyltransferase 10 (pp-GaNTase 10, EC 2.4.1.41) protein. Human pp-GaNTase 10 is located to chromosome 7q11.2. (I) can be used in gene therapy. Molecules of the present invention can be used in therapy, particularly to prevent or treat a disorder associated with decreased expression or activity of pp-GaNTase. The sequences given in ABV85011 to ABV86689 and ABP33502 to ABP53504 are given in the exemplification of the present invention. N.B. The sequence data for this patent is not represented in the printed specification but is based on sequence information supplied by the
                                                                                                                                                        Human, UDP-GalNAc:polypeptide N-acetylgalactosaminyltransferase 10;
pp-GaNTase 10; EC 2.4.1.41; chromosome 7q11.2; gene therapy; scanning;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2.8%; Score 12.8; DB 1; Length 17;
87.5%; Pred. No. 4e+02;
ative 0; Mismatches 2; Indels
                                                                                                                    Human pp-GaNTase 10 scanning 17-mer SEQ ID NO:739.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000670.
23-MAY-2001; 2001WS-00864761.
BP
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2001WO-US000664
                                                                                                                                                                                                                                                                                                                                                                                       2002EP-00001161
ABV85746 standard; DNA; 17
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                                                                            (first entry)
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                                                                                                                                                                                                                                             sapiens.
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                                                                              11-DEC-2002
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                                                                                                                                                                                                                                                                Synthetic.
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                                      ABV85746;
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Kim J;

Rice MC,

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The invention relates to an oligonucleotide for targeted alteration of a genetic sequence, which comprises a single-stranded oligonucleotide caving a DNA domain. The DNA domain has at least one mismatch with respect to the genetic sequence to be altered and further comprises chemical modification, an LNA modification, two or more consist of o-methyl modification, an LNA modification, two or more phosphorothicate linkages on a terminus, or a combination of any two or more of these modifications. The oligonucleotides are useful for conference of these modifications of plant genetic information. The oligonucleotides are particularly useful for creating plants with desired phenotypes, e.g. environmental or abjotic stress tolerance, improved cutritional value (e.g. altering amino acid content of plants with desired conferring amino acid over production), herbicide resistance (e.g. altering amino acid content of plants eresistance).

Conferring amino acid over production), herbicide resistance (e.g. altering amino acid over production), citazine resistance, conference modified oil production, modified starch production (e.g. increased starch or production of waxy starch), altered floral content (e.g. increased starch or production of waxy starch), altered floral content (e.g. reduced palmitate, increased stearte or reduced linolent acid).

Constitutional value of seaso useful for producting albino mutants for the altering of increases are sequence represents a genome of altering of increases.
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                                                                                                                    Claim 7; Page 87; 220pp; English.
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production.
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Gaps

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Indels

2.8%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 4e+02;

Pred. No. 4e+02; 0; Mismatches

Conservative

Best Local Similarity Matches 14; Conserv

Query Match

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Gaps

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343 GCGAGGTTCAGGCCTT 358

14; Conservative

Matches

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Local Similarity

ABK25391 standard; DNA; 17 BP.

RESULT 438

ABK25391

ABK25391 ID ABK2 XX AC ABK2

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Chromosomal genomic alteration; genome altering oligonucleotide; PCR; ss; o-methyl modification; LNA modification; phosphorothioate linkage; DNA repair; DNA alteration; environmental tolerance; hygromycin-B; abiotic stress tolerance; improved nutritional value; hygromycin; primer; amino acid over production; herbicide resistance; glyphosate resistance; pophyric herbicide resistance; sulphonylurea herbicide resistance; pophyric herbicide resistance; riazine resistance; disease resistance; modified oil production; modified starch production; waxy starch; modified fatty acid content; reduced plant; albino mutant; increased stearate production; reduced plantate production; albino plant; photosynthetic process.
                                                                                                                                  Male-sterile plant producing genome altering oligonucleotide #292.
                                                                       ABK25392 standard; DNA; 17 BP
                    creaccreaceccer 16
                                                                                                                (first entry)
410 CTGAGCTGTGGGACGT
                                                                                                                                                                                                                                                                                                                          WO200192512-A2.
                                                                                                                 09-APR-2002
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                                                   RESULT 439
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Kim J; Rice MC, 01-JUN-2000; 2000US-0208538P. 30-OCT-2000; 2000US-0244989P. 27-MAR-2001; 2001US-00818875. 01-JUN-2001; 2001WO-US017672. Gamper HB, (UYDE) UNIV DELAWARE. Kmiec EB,

New oligonucleotides with modified nuclease-resistant termini, useful for creating plants with desired phenotypes, e.g. stress tolerance, improved nutritional value, herbicide or disease resistance, or modified oil production.

WPI; 2002-106307/14.

Claim 7; Page 87; 220pp; English.

The invention relates to an oligonucleotide for targeted alteration of a genetic sequence, which comprises a single-stranded oligonucleotide CC flaving a DNA domain. The DNA domain has at least one mismatch with respect to the genetic sequence to be altered and further comprises chemical modifications of the oligonucleotide. The chemical modifications of the oligonucleotide. The chemical modifications of the oligonucleotide are useful for consist of o-methyl modification, an LNA modification, two or more phosphorothioate linkages on a terminus, or a combination of any two or more of these modifications. The oligonucleotides are useful for directing repair or alteration of plant genetic information. The oligonucleotides are particularly useful for creating plants with desired phenotypes, e.g. environmental or abiotic stress tolerance, improved conferring amino acid over production), herbicide resistance (e.g. altering amino acid over production), herbicide resistance (e.g. glyphosate resistance, imidazolinome and sulphonylurea herbicide, cresistance, porphyric herbicide resistance or triazine resistance), disease resistance, modified oil production, modified starch production (e.g. increased starch or production of waxy starch), altered floral morphology (e.g. male-sterile plants) or modified fatty acid content (e.g. reduced palmitate, increased stearate or reduced linolenic acid).

The invention relates to a nucleic acid molecule (I) which down regulates expression of an Ets-related gene (ERG). (I) is useful for treating conditions selected from cancer, lymphoma, Eaving's sarcoma, melanoma, tumour angiogenesis, diabetic retinopathy, macular degeneration, neovascular glaucoma, myopic degeneration, arthritis, psoriasis, vertuca valgaris, angiofibroma of tuberous sclerosis, port-whom stains, Sturge Weber syndrome, Kippel-Trenaunay-Weber syndrome, Osler-Weber-rendu syndrome, leukaemia, osteoporosis and wound healing. (I) is useful for treatment. The method comptises the use of one or more therapies the treatment. The method comprises the use of one or more therapies the treatment. Leukaemia or tumour angiogenesis is treated by administering (I) to the patient in angiogenesis is treated by administering (I) to the patient in conjunction with one or more of other therapies such as radiation or chemotherapy treatment. (I) is useful for reducing ERG activity in a

Novel polynucleotide which down regulates expression of Ets-related gen useful for treating cancer, diabetic retinopathy, macular degeneration, arthritis, psoriasis, verruca vulgaris and Sturge Weber syndrome.

Claim 4; Page 74; 149pp; English.

Mclaughlin F, Randi AM;

Mcswiggen JA,

Jarvis T, Von Carlowitz I,

WPI; 2002-082995/11.

(RIBO-) RIBOZYME PHARM INC.

(GLAX) GLAXO GROUP LTD.

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The oligonucleotides are also useful for producing albino mutants for the analysis of photosynthetic processes. This sequence represents a genome altering oligonucleotide of the invention
                                                                                                                                                                                                                                                                                                         Human; hammerhead ribozyme; cytostatic; antitumour; antidiabetic; ophthalmological; antiarthritic; antipsoriatic; virucide; osteopathic; vulnerary; cancer; lymphoma; Ewing's sarcoma; melanoma; psoriasis; tumour angiogenesis; diabetic retinopathy; macular degeneration; neovascular glaucoma; myopic degeneration; arthritis; verruca vulgaris; angiofibroma of tuberous sclerosis; port-wine stain; wound healing; Sturge Weber syndrome; Kippel-Trenaunay-Weber syndrome; leukaemia; ss; oster-Weber-rendu syndrome, leukaemia; osteoporosis; DNAzyme; inozyme;
                                                                                                      Gaps
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0
                                                                                                                                                                                                                                                                                  Human BRG hammerhead ribozyme target sequence, Seq ID No 839.
                                                                          2.8%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 4e+02;
                                                                                                     Indels
                                                                                                    7
                                                Sequence 17 BP; 3 A; 7 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                    0; Mismatches
                                                                                                                           425
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                                                                                                                        410 CTGAGCTGTGGGACGT
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                                                                    Query Match
Best Local Similarity
Matches 14; Conserv
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ERG gene, by contacting the cell with (I). (I) is useful for cleaving RNA of ERG gene, by contacting (I) with RNA, in the presence of a divalent cation such as Mg2+. (I) is useful for diagnosis of conditions and diseases related to the expression of ERG, and as diagnostic tool to examine genetic drift and mutations within diseased cells or to detect the presence of ERG RNA in a cell. (I) is useful for specifically targeting genes that share homology with ERG gene or ERG fusion genes. ABK17354-ABK22719 represent nucleic acids, including antisense and enzymatic nucleic acid molecules which regulate expression of ERG, and related PCR primers of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel polynucleotide which down regulates expression of Ets-related gene, useful for treating cancer, diabetic retinopathy, macular degeneration, arthritis, psoriasis, verruca vulgaris and Sturge Weber syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; hammerhead ribozyme; cytostatic; antitumour; antidiabetic; ophthalmological; antiarthritic; antipsoriatic; virucide; osteopathic; vulnerary; cancer; lymphoma; Ewing's sarcoma; melanoma; psoriasis; tumour angiogenesis; diabetic retinopathy; macular degeneration; neovascular glaucoma; myopic degeneration; arthritis; verruca vulgaris; angiofibroma of tuberous sclerosis; port-wine stain; wound healing; Sturge Weber syndrome; Kippel-Trenaunay-Weber syndrome; leukaemia; 88;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human ERG hammerhead ribozyme target sequence, Seq ID No 838
                                                                                                                                                                                                                                                                                      2.8%; Score 12.8; DB 1; Length 17;
87.5%; Pred. No. 4e+02;
ive 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABK18191 standard; RNA; 17 BP.
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                                                                                                                                                                                                                                                                                                                                                                               7 CGGAGGGTGGGCCTGG
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Matches 14; Conservative
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ID ABK1
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treating a patient having a condition associated with the level of ERG, by contacting cells of the patient with (1) under conditions suitable for the treatment. The method comprises the use of one or more therapies under conditions suitable for the treatment. Leukaemia or tumour angiogenesis is treated by administering (1) to the patient in conjunction with one or more of other therapies such as radiation or chemotherapy treatment. (1) is useful for reducing ERG activity in a cell, by contacting the cell with (1). (1) is useful for cleaving RNA of ERG gene, by contacting (1) with RNA, in the presence of a divalent cation such as Mg2+. (1) is useful for diagnosis of conditions and diseases related to the expression of ERG, and as diagnosit tool to examine genetic drift and mutations within diseased cells or to detect the presence of ERG RNA in a cell. (1) is useful for specifically targeting genes that share homology with ERG gene or ERG fusion genes. ARXI314-ABK22719 represent nucleic acide, including antisense and enzymatic nucleic acid molecules which regulate expression of ERG, and carginate expression of ERG, and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human, POSHL 1; SH3 domain, POSH-like signalling protein 1; oncogene; Rho GTPase; signal transduction; gene expression; cancer; vaccine;
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                                                                                                                                                                                                                                                                                                                                                 2.8%; Score 12.8; DB 1; Length 17;
87.5%; Pred. No. 4e+02;
tive 0; Mismatches 2; Indel8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human POSHL1 scanning oligonucleotide SEQ ID NO 716.
                                                                                                                                                                                                                                                                                                                 Sequence 17 BP; 2 A; 12 C; 2 G; 0 T; 1 U; 0 Other;
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2001WO-US000668.
2001WO-US000669.
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30-JAN-2001; 2
30-JAN-2001; 2
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ID ABV9
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The invention relates to an isolated SH3 domain (POSH)-like signalling protein 1 (POSH1.1) polypeptide (I), comprising a sequence of 730 amino acids (SI, ABB8399), a sequence having 65$ sequence identity to (SI), (SI) having 95$ deviations, especially conservative substitutions or a fragment of the sequences comprising at least 8 contiguous amino acids. Human POSH1 is a proteo-oncogene/oncogene product that functions as an adaptor protein that interacts with Rho family small GTPases as well as downstream components of the signal transduction pathway. (I) is useful for for identifying a specific binding partner. (I) and mucleic acids (II) encoding (I) are useful for diagnosing, monitoring disease and traating caused by altered expression of human POSH11 including diagnosing and treating cancer, they useful in the development of vaccines and (II) is useful in gene therapy. (II) is useful for constructing microarrays which are useful for measuring and for surveying gene expression and creating transgenic non-human animals capable of producing the proteins. The present sequence is that of a scanning oligomorleotide useful in examples of the invention. Note: The present sequence did not form part of the printed specification, but is based on sequence information supplied to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human, POSHL 1; SH3 domain, POSH-like signalling protein 1, oncogene, Rho GTPase; signal transduction, gene expression, cancer, vaccine,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human POSHL1 scanning oligonucleotide SEQ ID NO 715.
                                                                                                                                                                                                                                                                                                                                                                                               Sequence 17 BP; 3 A; 6 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                          Derwent by the European Patent Office
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30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000667.
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2001WO-US000669.
2001WO-US000670.
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2001US-0328205P
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2001WO-US000664
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABV90002;
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protein 1 (POSHL 1) polypeptide (1), comprising a sequence of 730 amino acids (S1, ABB31999), a sequence having 65% sequence identity to (S1), acids (S1) ABB31999), a sequence having 65% sequence identity to (S1), acids (S1) having 95% deviations, especially conservative substitutions or a fragment of the sequences comprising at least 8 contribuous amino acids. Human POSHL 1 is a proto-oncogene product that functions as an adaptor protein that interacts with Rho family small GrPases as well as daptor protein that interacts with Rho family small GrPases as well as daptor protein that interacts with Rho family small GrPases as well as considered components of the signal transduction pathway. (I) is useful for or identifying a specific binding partner. (I) and nucleic acids (II) caused by altered expression of human POSHL1 including disease and treating caused by altered expression of human POSHL1 including diagnosing and treating causeful for constructing microarrays which are useful for measuring and for surveying gene expression and creating transgenic non-human animals capable of producing the proteins. The present sequence is that of a scanning oligomore of the invention. Note: The present sequence did not form part of the printed specification, but is based on sequence information supplied to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a typing kit for judging human leukocyte antigen (HLA) genotype of a sample by hybridising a substrate on which 10-24 base
                                                                                                  The invention relates to an isolated SH3 domain (POSH)-like signalling nrotein 1 (POSHL 1) polypeptide (I), comprising a sequence of 730 amino comprising a sequence of 730 amino comprising a sequence of 730 amino compression (POSHL 1) polypeptide (I), comprising a sequence of 730 amino compression (POSHL 1) polypeptide (I), comprising a sequence of 730 amino compression (POSHL 1) polypeptide (I), compress
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immunogenetic, transplantation, genetic disease, ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 2.8%; Score 12.8; DB 1; Length 17; Best Local Similarity 87.5%; Pred. No. 4e+02; Matches 14; Conservative 0; Mismatches 2; Indels
                                                                              Example 2; SEQ ID NO 715; 60pp + Sequence Listing; English.
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activity of human POSHL1.
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oligonucleotides (ABL30512-ABL31809) originating in the sequences of genes e.g. belonging to HLA class I antigens on human genome and containing gene polymorphisms as alloantigens have been immobilised as primers for amplification of cleaved nucleic acids relating to gene polymorphisms. The method is useful for judging HLA genotypes of individuals by determining immunogenetic differences before transplanting between them, providing genetic information to decide compatibility of organ and tissue for transplantation e.g. of bone marrow, kidney, liver, pancreas, Langerhans islet in pancreas and cornea, susceptibility diagnosis of genetic diseases and identifying individuals
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to nucleic acid molecules that modulate replication of the West Nile Virus (WNV). The nucleic acid molecules are useful for treating a condition related to WNV infection e.g. pancreatitis, encephalitis, myocarditis, meningitis, neurologic infection, hepatitis, liver failure, hepatocellular carcinoma or cirrhosis. The nucleic acid molecule is selected from the group of ribozymes consisting of Hammerhead, Inozyme, G-claaver, DNAzyme, Amberzyme and Zinzyme. The nucleic acid molecules further comprise at least five ribose residues, at least ten 2-0-methyl modifications, phosphorothicate linkages on at least three of the 5' terminal nucleocides and a 3' end modification of a 3' inverted abasic moiety. Nucleic acid molecules SEQ ID NO 1 to 37080
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                                                                                                                                                                                                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                                        2.8%; Score 12.8; DB 1; Length 17;
87.5%; Pred. No. 4e+02;
ive 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                  Sequence 17 BP; 1 A; 6 C; 7 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                       379 CGGAGCGAGTCCCCGC 394
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nes 14; Conservative
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(MCSW/) MCSWIGGEN J A.
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encephalitis, myocarditis, meningits, neurologic infection, hepatitis, liver failure, hepatocallular carcinoma or cirrhosis. The nucleic acid molecule is selected from the group of ribozymes consisting of mammerhead, Inozyme, G-cleaver, DNAzyme, Amberzyme and Zinzyme. The nucleic acid molecules further comprise at least tive ribose residues, at least ten 2'-O-methyl modifications, phosphorothioate linkages on at least three of the 5' terminal nucleotides and a 3' end modification of a 3'-3' inverted abasic moiety. Nucleic acid molecules SEQ ID NO 1 to 37080 are claimed; however, SEQ ID NO 2194-2206 and 17502-17514 are not given in the specification. The present sequence is that of a nucleic acid
are claimed; however, SEQ ID NO 2194-2206 and 17502-17514 are not given in the specification. The present sequence is that of a nucleic acid molecule of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New nucleic acid molecule that modulates replication of West Nile Virus (WNV), useful for treating a condition related to WNV infection e.g. pancreatitis, meningitis, hepatocellular carcinoma or cirrhosis.
                                                                                                                                                     Сарв
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WNV; West Nile Virus; antiinflammatory; cytostatic; hepatotropic; virucide; neuroprotective; antibacterial; replication; panoreatitis; encephalitis; myocarditis; minfection; hepatitis; liver failure; cancer; cirrhosis; Hammerhead; Inozyme; DNAzyme;
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                                                                            G; 0 T; 5 U; 0 Other;
                                                                                                             2.8%; Score 12.8; DB 1
52.5%; Pred. No. 4e+02;
ve 4; Mismatches
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1 UGAGCCGCAGGUCUCU 16
                                                                            Sequence 17 BP; 2 A; 5 C; 5
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                                                                                                                                Local Similarity 62.
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DB 1; Length 17;

2.8%; Score 12.8;

Query Match

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to nucleic acid molecules that modulate replication
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New nucleic acid molecule that modulates replication of West Nile Virus (WNV), useful for treating a condition related to WNV infection e.g. pancreatitis, meningitis, hepatocellular carcinoma or cirrhosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WNV, West Nile Virus, antiinflammatory; cytostatic; hepatotropic; virucide, neuroprotective; antibacterial; replication; panoreatitis; encephalitis; myocarditis; meningitis; infection; hepatitis; liver failure; cancer; cirrhosis; Hammerhead; Inozyme; DNAzyme;
                                    Gaps
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                              2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WNV minus strand Amberzyme substrate SEQ ID NO 15002.
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   87.5%; Pred. No. 4e+02;
                                 Mismatches
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                        0;
                                                                                                                                                                                                                                                                                        ACN14999 standard; RNA; 17 BP.
                                                                                    316 TCAGCCGCGGGTCTCT 331
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-OCT-2001; 2001WO-US048350.
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                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
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                              14; Conservative
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(MCSW/) MCSWIGGEN J A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Amberzyme; Zinzyme; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2002-706994/76.
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Best Local Similarity
Best Local Similarity
Matches 14; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   West Nile Virus.
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                                                                                                                                                                                                                                                                                                                                                 ACN14999;
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                                                                                                                                                                                                                                     447
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New nucleic acid molecule that modulates replication of West Nile Virus

20-OCT-2000; 2000US-0242411P.

RIBO-) RIBOZYME PHARM INC

(MCSW/) MCSWIGGEN J A.

BLAT/)

Mcswiggen JA;

Blatt L,

WPI; 2002-706994/76.

19-OCT-2001; 2001WO-US048350.

06-SEP-2002

West Nile Virus. WO200268637-A2.

WNV; West Nile Virus; antiinflammatory; cytostatic; hepatotropic; virucide; neuroprotective; antibacterial; replication; pancreatitis; encephalitis; myocarditis; meningitis; infection; hepatitis; liver failure; cancer; cirrhosis; Hammerhead; Inozyme; DNAzyme; Amberzyme; Zinzyme; ss.

WNV Hammerhead Ribozyme substrate SEQ ID NO 405.

(first entry)

22-APR-2004

ACN00415;

ACN00415 standard; RNA; 17 BP.

RESULT 448

ACN00415,

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (WNV), useful for treating a condition related to WNV infection e.g. pancreatitis, meningitis, hepatocellular carcinoma or cirrhosis.
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Pred. No. 4e+02;
0; Mismatches 2; Indels
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ID ACN03272 standard; RNA; 17 x

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AC ACN03272;

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2.8
Best Local Similarity 87.5
Matches 14; Conservative
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Gaps

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TCTCGCTGACTTTCAG 118

103

Conservative

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liver failure; cancer; cirrhosis; Hammerhead; Inozyme; DNAzyme;
             Amberzyme; Zinzyme; ss
                                                                                                                                                                                                                                              WPI; 2002-706994/76.
                                     West Nile Virus.
                                                              #0200268637-A2
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                                                                                       06-SEP-2002,
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                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to nucleic acid molecules that modulate replication of the West Nile Virus (WNV). The nucleic acid molecules are useful for treating a condition related to WNV infection e.g. pancreatitis.

encephalitis, myozaccalitis, meningnitis, neurologic infection, hepatitis, liver failure, hepatocellular carcinoma or cirrhosis. The nucleic acid molecule is selected from the group of ribozymes consisting of Hammerhead, Inozyme, G-cleaver, DMAZyme, Amberzyme and Zinzyme. The nucleic acid molecules further comprise at least five ribose residues, least three of the S' terminal nucleotides and a 3' end modification of a 3'-3' inverted abasic moiety. Nucleic acid molecules SEQ ID NO 1 to 37080 are claimed; however, SEQ ID NO 2194-2206 and 17502-17514 are not given in the specification. The present sequence is that of a nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                New nucleic acid molecule that modulates replication of West Nile Virus (WNV), useful for treating a condition related to WNV infection e.g. pancreatitis, meningitis, hepatocellular carcinoma or cirrhosis.
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                                                WNV, West Nile Virus, antiinflammatory; cytostatic; hepatotropic; viruclé; neuroprotective; antibacterial; replication; pancreatitis; encephalltis; myocarditis; meningitis; infection; hepatitis; liver failure; cancer; cirrhosis; Hammerhead; Inozyme; DNAzyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WNV; West Nile Virus; antiinflammatory; cytostatic; hepatotropic; virucide; neuroprotective; antibacterial; replication; pancreatitis; encephalitis; myocarditis; meningitis; infection; hepatitis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WNV minus strand DNAzyme substrate SEQ ID NO 14013.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 17 BP; 5 A; 5 C; 5 G; 0 T; 2 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 23; SEQ ID NO 3275; 495pp; English.
                         WNV Inozyme substrate SEQ ID NO 3275.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17 TGAGCCGCAGGTCTCT 2
                                                                                                                                                                                                                                                          (RIBO-) RIBOZYME PHARM INC.
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(first entry)
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                                                                                                                                                                                                                                                                                                              Blatt L, Mcswiggen JA;
                                                                                                      Amberzyme; Zinzyme; ss
                                                                                                                                                                                                                                                                                    (MCSW/) MCSWIGGEN J A.
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                                                                                                                             West Nile Virus.
                                                                                                                                                     WO200268637-A2
 22-APR-2004
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of the West Nile Virus (WNV). The nucleic acid molecules are useful for treating a condition related to WNV infection e.g. pancreatitis, encephalitis, meningits, neurolegic infection, hepatitis, meningits, neurolegic infection, hepatitis, meningits, neurolegic infection, hepatitis, molecule is selected from the group of ribozymes consisting of Hammerhead, Inozyme, G-Cleaver, DNAzyme, Amberzyme and Zinzyme. The nucleic acid molecules further comprise at least five ribose residues, at least the 2-O-methyl modifications, phosphorochioate linkages on at least three of the 5' terminal nucleotides and a 3' end modification of a 3'-3' inverted abasic moiety. Nucleic acid molecules SEQ ID No 1 to 37080 are claimed, however, SEQ ID No 2194-2206 and 17502-17514 are not given in the specification. The present sequence is that of a nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to nucleic acid molecules that modulate replication
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New nucleic acid molecule that modulates replication of West Nile Virus (WNV), useful for treating a condition related to WNV infection e.g. pancreatitis, meningitis, hepatocellular carcinoma or cirrhosis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 23; SEQ ID NO 14013; 495pp; English
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19-OCT-2001; 2001WO-US048350.
                                                                        20-OCT-2000; 2000US-0242411P
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                                                                                                                                                (RIBO-) RIBOZYME PHARM INC.
(BLAT/) BLATT L.
(MCSW/) MCSWIGGEN J A.
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US09436060A-16.rng.sl

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Best Loca
Matches
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                                                                                                                                                                                                                                                                                 The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is cancoded at chromosome 7422.1, MDZ4 is encoded at chromosome 6721.3-22.2, MDZ7 is encoded at chromosome 6721.3-22.2, MDZ7 is encoded at chromosome 1601.2 and MDZ12 is encoded at chromosome 572.1, mDZ7 is encoded at chromosome 1601.2 and MDZ12 is encoded at chromosome 1601.2 and MDZ12 and MDZ12 is encoded at chromosome 1602.7 and MDZ12 and MDZ12 and MDZ12 and useful in therapy, or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12 genetic locus. The probes are useful as therapeutic agents for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ7; MDZ1; chromosome 7g22.1; chromosome 6p21.3-22.2; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss.
                                                                                                                                                                       New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7 or MDZ12, e.g. cancer.
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                                                                                                                                                                                                                                                        Example 8; SEQ ID NO 815; 103pp; English.
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              30-JUL-2002; 2002EP-00016874.
                                            02-AUG-2001; 2001US-00922181
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Best Local Similarity 87.5%;
Matches 14; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25 GGGGTGGTGGCCATTT 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GGGGTTGGGGCCATTT 16
                                                                                                           Gu Y, Nguyen C;
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                                                                                                                                           WPI; 2003-423107/40.
                                                                            (AEOM-) AEOMICA INC
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                                                                                                           Shannon M,
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                        New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7 or MDZ12, e.g. cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; ribozyme; short interfering RNA; siRNA; HER2; K-Ras; enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV; anti-rheumatic; cancer; AIDS; ss.
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                                                                                                                                                                                                                                                                                                                                  Sequence 17 BP; 2 A; 3 C; 8 G; 4 T; 0 U; 0 Other;
                                                                                          Example 8; SEQ ID NO 813; 103pp; English.
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es 14; Conservative
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WPI; 2003-423107/40.
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ABZ66530
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                                                                 acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HER2, K-Ras, H-Ras, N-Ras, human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-rheumatic activity. The nucleic acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung prostate, bladder, or pancreatic cancer, and HIV infection, and AIDS. The sequences shown in ABE25889 - ABE26216, ABE265531, ABE26520 - ABE26524, ABE26530 - ABE26520 - ABE26524, and ribozymes of the invention
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                                                The invention relates to a novel short interfering RNA (siRNA) nucleic
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enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV;
anti-rheumatic; cancer; AIDS; ss.
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  Claim 58; Page 129; 185pp; English
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06-JUN-2001; 2001US-0296249P.
10-SEP-2001; 2001US-0318471P.
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Matches 14; Conservative
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shown in ABZ59889 - ABZ65216, ABZ64544 - ABZ65531, ABZ66520 - ABZ66524,
ABZ66530 - ABZ66585 represent substrate/target sequences for the human
ribozymes of the invention
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                                                                                                                                                                                        2.8%; Score 12.8; DB 1; Length 17; 77.5%; Pred. No. 4e+02; ve 0; Mismatches 2; Indels
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                                                                                                                                  Seguence 17 BP; 2 A; 11 C; 1 G; 0 T; 3 U; 0 Other;
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06-JUN-2001; 2001US-0296249P.
10-SEP-2001; 2001US-0318471P.
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Best Local Similarity 87.5%
Local 14; Conservative
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hes 14; Conservative
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Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences.
                                               Human; ribozyme; short interfering RNA; siRNA; HER2; K-Ras;
enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV;
anti-rheumatic; cancer; AIDS; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 58; Page 114; 185pp; English
                    Human H-Ras DNAzyme target #179,
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10-SEP-2001; 2001US-0318471P.
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                                                                                                                                    WO200297114-A2
                                                                                                         Homo sapiens
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                                                                                                                                                                                                Human, ribozyme; short interfering RNA; siRNA; HER2; K-Ras;
enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV;
anti-rheumatic; cancer; AIDS; ss.
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                                                                                                                                                                       Human HER2 DNAzyme substrate #20.
                                                                                   ABZ64563 standard; RNA; 17 BP
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              16 GCCGCGGTCGGGCGGG 1
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les 14; Conservative
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The invention relates to a novel short interfering RNA (siRNA) nucleic acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HER2, K.Ras, H.Ras, N.Ras, human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-HER2, K.Ras, H.Ras, and HIV activity in a cell. The nucleic acid stream the same useful for treating breast, ovarian, colorectal, lung, prostate, bladder, or pancreatic cancer, and HIV infection, and AIDS. The sequences shown in ABES9899 - ABES6216, ABE65531, ABE66520 - ABE66524, riboszymes of the invention
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Matches
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Gaps

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21 GGGAGGGTGGTGGCC 36

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Matches

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ABZ61388 standard; RNA; 17

RESULT 457

ABZ61388

ABZ61388;

21-MAR-2003 (first entry)

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Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus
                                     Nucleic acid molecule, Hepatitis C virus; HCV; Hepatitis B virus; HBV; RNA stability; RNA expression; RNA synthesis; antisense; enzymatic nucleic acid; hammerhead ribozyme; DNAzyme; inozyme; zinzyme;
                                                                            amberzyme; G-cleaver ribozyme; decoy molecule; aptamer;
HBV reverse transcriptaes; Enhancer I region; viral replication;
degenerative; disease state; HBV infection; HCV infection; cirrhosis;
liver failure; hepatocellular carcinoma; hepatotropic; cytostatic;
             minus strand DNAzyme substrate sequence #1478.
                                                                                                                                 virucide; antiinflammatory; substrate; ss.
                                                                                                                                                                                                                                                                 26-MAR-2001; 2001US-00817879.
08-JUN-2001; 2001US-00877478.
08-JUN-2001; 2001US-0296876P.
24-OCT-2001; 2001US-0335059P.
05-DEC-2001; 2001US-0337055P.
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                                                                                                                                                           Hepatitis C virus
                                                                                                                                                                                      WO200281494-A1
                                                                                                                                                                                                               17-0CT-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                infection.
                                                                                                                                                                                                                                                                                                                                                                                                                    (PAVC/)
(LEEP/)
                                                                                                                                                                                                                                                                                                                                                 (RIBO-)
                                                                                                                                                                                                                                                                                                                                                                                                                                               (DRAP/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                          (ROBE/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Draper
                                                                                                                                                                                                                                                                                                                                                               (BLAT/
                                                                                                                                                                                                                                                                                                                                                                                          MCSW/
                                                                                                                                                                                                                                                                                                                                                                                                      (MORR/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Blatt
 The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HCV) or and enzymatic nucleic acid molecules include antisense and enzymatic nucleic acids such as hammerhead ribozymes, DNAzymes, inozymes, zinzymes, amberzymes, and G-cleaver ribozymes, DNAzymes, or nozymes, zinzymes, amberzymes, and G-cleaver ribozymes. Also disclosed transcriptase and/or HBV reverse transcriptase primer sequences, as well as oligonucleotides that specifically bind the Enhancer I region of HBV genes and HBV viral replication. Also disclosed is a method for screening compounds and/or potential therapies directed against HBV, and compounds that modulate the expression and/or halso disclosed is a method for screening compounds and/or potential therapies directed against HBV, and compounds methods of the invention are useful for the treatment of degenerative and disease states related to HBV and HCV infection, replication and gene carcinoma. The present sequence represents a substrate for one of the HCV binazyme or minus strand DNAzyme sequences disclosed in the present
                                                                                                                                                                                                                                                                                                                                                                                                      Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus
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                                                                                                                                                                                                                                                                                                                                      Lee
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  / Match 2.8%; Score 12.8; DB 1; Length 17; Local Similarity 75.0%; Pred. No. 4e+02; tes 12; Conservative 2; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                     Pavco P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 17 BP; 1 A; 1 C; 11 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                     Morrissey D,
                                                                                                                                                                                                                                                                                                                                     Mcswiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; Page 259; 387pp; English
                                                                                                                                 2001US-00877478.
2001US-0296876P.
                                                                                                                                                          24-OCT-2001; 2001US-0335059P.
                                                                                          26-MAR-2002; 2002WO-US009187
                                                                                                                                                                                                  RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                     Macejak D,
Roberts E;
                                                                                                                                                                                                                           MACEJAK D.
MCSWIGGEN J.
MORRISSEY D.
                                                                                                                                                                                                                                                                                                                                                                            WPI; 2003-229207/22.
                                                                                                                                                                                                                                                                               LEE P.
DRAPER K.
ROBERTS E.
                                                                                                                                                                                                                                                                                                                                   Macejak
            Hepatitis C virus
                                                                                                                                                                                                                                                                     PAVCO P.
                                      WO200281494-A1
                                                                                                                                                                                                                BLATT
                                                                                                                                 08-JUN-2001;
                                                                                                                    26-MAR-2001;
                                                                 17-OCT-2002
                                                                                                                                                                                                                                                                                                                                     Blatt L, M
Draper K,
                                                                                                                                                                                                                                                                                                                                                                                                                                 infection.
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                                                                                                                                                                                                                                                    (MORR/) 1
(PAVC/) 1
(LEEP/) 1
(DRAP/) 1
(ROBE/) 1
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                                                                                                                                                                                                 (RIBO-)
(BLAT/)
(MACE/)
(MCSW/)
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RIBOZYME PHARM INC

MACEJAK D. MCSWIGGEN J. MORRISSEY D.

ď Lee

o, Pavco

Morrissey D,

Mcswiggen J,

Macejak D, Roberts E;

ROBERTS E DRAPER K. PAVCO P.

LEE

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The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HCV) or Hepatitis B virus (HBV) RNA. The nucleic acid molecules include antisense and enzymatic nucleic acids such as hammerhead ribozymes, DNAzymes, amberzymes, amberzymes, and G-cleaver ribozymes, Also disclosed are nucleic acid decoy molecules and G-cleaver ribozymes. Also disclosed transcriptase and/or HBV reverse transcriptase primer sequences, as well as oligomucleotides that specifically bind the Enhancer I region of HBV genes and HBV viral replication. Also disclosed is a method for screening compounds and/or potential therapies directed against HBV, and compounds compounds and/or potential therapies directed against HBV, and compounds methods of the invention are useful for the treatment of degenerative and disease states related to HBV and HCV infection, replication and gene archivesion such as cirrhosis, liver failure, and hepatocellular carcinoma. The present sequence represente a substrate for one of the HCV DNAzyme or minus strand DNAzyme sequences disclosed in the present
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 17 BP; 1 A; 6 C; 6 G; 0 T; 4 U; 0 Other;
Claim 1; Page 301; 387pp; English.
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Best Local Similarity 87.55
Matches 14; Conservative
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Gaps

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35

20 TGGGAGGGGTGGTGGC

ઠ 8

Matches

ACD64287 standard; RNA; 17

RESULT 459

(first entry)

30-SEP-2003

ACD64287;

ACD64287/c ID ACD64 XX AC ACD64 XX DT 30-SE

US09436060A-16.rng.sl

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The present invention relates to nucleic acid molecules which modulate
the synthesis, expression and/or stability of Hepatitis C virus (HCV) or
Hepatitis B virus (HBV) RNA. The nucleic acid molecules include antisense
and enzymatic nucleic acids such as hammerhead ribozymes. DNAzymes,
c inozymes, zinzymes, amberzymes, and G-cleaver ribozymes. Also disclosed
are nucleic acid decoy molecules and aptamers that bind to HBV reverse
transcriptase and/or HBV reverse transcriptase primer sequences, as well
as oligonucleotides that specifically bind the Enhancer I region of HBV
C as oligonucleotides that specifically bind the Enhancer I region of HBV
C as oligonucleotides that specifically bind the Enhancer I region of HBV
C as oligonucleotides that specifically bind the Enhancer I region of HBV
C and of the invention are used to modulate the expression of HBV
c hat modulate the expression and/or replication of HCV. The compounds
methods of the invention are useful for the treatment of degenerative and
disease states related to HBV and HCV infection, replication and gene
expression such as cirrhosis, liver failure, and hepatocellular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus
                                                                                                                                                                                                                                           Nucleic acid molecule; Hepatitis C virus; HCV; Hepatitis B virus; HBV; RNA stability; RNA expression; RNA synthesis; antisense; enzymatic nucleic acid; hammerhead ribozyme; DNAzyme; inozyme; zinzyme; amberzyme; G-cleaver ribozyme; decoy molecule; aptamer; HBV reverse transcriptase; Enhancer I region; viral replication; degenerative; disease state; HBV infection; HCV infection; cirrhosis; liver failure; hepatocellular carcinoma; hepatotropic; cytostatic; virucide; antiinflammatory; substrate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Lee P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mcswiggen J, Morrissey D, Pavco P,
                                                                                                                                                                                                          HCV minus strand DNAzyme substrate sequence #1362.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1; Page 299; 387pp; English.
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08-UUN-2001; 2001US-00877478.
08-UUN-2001; 2001US-0296876P.
24-OCT-2001; 2001US-0337055P.
05-DEC-2001; 2001US-0337055P.
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                     (7
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                                                                                                      ACD64059 standard; RNA; 17
(first entry)
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Roberts E;
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MCSWIGGEN J.
MORRISSEY D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-229207/22.
                                                                                                                                                                                                                                                                                                                                                                                                      Hepatitis C virus.
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DRAPER K.
ROBERTS E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PAVCO P.
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Draper K,
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                                                                                                                                          ACD64059;
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(BLAT/)
(MACE/)
(MCSW/)
(MORR/)
(PAVC/)
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(ROBE/)
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carcinoma. The present sequence represents a substrate for one of the HCV DNAzyme or minus strand DNAzyme sequences disclosed in the present invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus
                                                                                                                                                                                                                                                                                                                                                         Nucleic acid molecule, Hepatitis C virus; HCV; Hepatitis B virus; HBV; RNA stability; RNA expression; RNA synthesis; antisense; enzymatic nucleic acid, hammerhead ribozyme; DNAzyme; inozyme; zinzyme; amberzyme; G-claever ribozyme; decoy molecule; aptemer; HBV reverse transcriptase; Enhancer I region; viral replication; degenerative; disease state; HBV infection; HCV infection; cirrhosis; liver failure; hepatocellular carcinoma; hepatotropic; cytostatic; virucide; antiinflammatory; substrate; ss.
                                                                                                                               Gaps
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                                                                                              2.8%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 4e+02; ative 0; Mismatches 2; Indels
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                                                                     Sequence 17 BP; 3 A; 6 C; 7 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Morrissey D,
                                                                                                                                                                                                                                                                                                                                HBV inozyme substrate sequence #352.
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                                                                                                                                                                                                                                                   BP.
                                                                                                                                                     208 GGACCTGCGGCGGGTC 223
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08-JUN-2001; 2001US-00877478.
08-JUN-2001; 2001US-0296876P.
24-CCT-2001; 2001US-0335059P.
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                                                                                                                                                                                                                                                  ACD52324 standard; RNA; 17
                                                                                                                                                                      GCACCTGCGGCGGCTC
                                                                                                                                                                                                                                                                                                       (first entry)
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                                                                                                                           Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BLATT L.
MACEJAK D.
MCSWIGGEN J.
MORRISSEY D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Macejak D,
                                                                                                            Local Similarity
les 14; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2003-229207/22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hepatitis B virus.
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LEE P.
DRAPER K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ROBERTS E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200281494-A1.
                                                                                                                                                                                                                                                                                                       24-SEP-2003
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Draper K,
                                                                                                                                                                             16
                                                                                                                                                                                                                                                                             ACD52324;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    infection.
                                                                                              Query Match
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(MACE/)
(MCSW/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (PAVC/)
(LEEP/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (RIBO-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ROBE/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (MORR/)
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                                                                                                                         Matches
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inozymes, zinzymes, amberzymes, and G-cleaver ribozymes. Also disclosed are nucleic acid decoy molecules and aptamers that bind to HBV reverse transcriptase and/or HBV reverse transcriptase primer sequences, as well as oligonuclectides that specifically bind the Enhancer I region of HBV genes and HBV viral replication. Also disclosed is a method for screening compounds and/or potential therapies directed against HBV, and compounds that modulate the expression and/or replication of HV. The compounds of the invention are useful for the treatment of degenerative and disease states related to HBV and HCV infection, replication and gen expression such as cirrhosis, liver failure, and hepatocellular carcinoma. The present sequence represents a substrate for one of the HBV ribozyme, inozyme, G-cleaver, zinzyme, DNAzyme or amberzyme sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to human phenotype-associated (PA) genes (I ; Appl3307-AbF13386) which contain a Single Nuclectide Polymorphism (SNP). The SNP is given in the sequence as a variant nucleotide. Also claimed are methods for screening for agents which regulate the activity of a PA gene and reagents that modulate the activity of a PA purclectide where the reagent is identified by the screening methods. The methods and compositions of the present invention are useful for prognosticating, diagnosing and treating cardiovascular diseases. Mypercential, hypertension, restenosis, arterial inflammation, used in the examples from the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cardiant; antiarteriosclerotic; vasotropic; cerebroprotective;
hypotensive; gene therapy; human; SNX9; Sorting Nexin 9; PCR; primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                                                                                                 2.8%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 4e+02; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SNX9 (Sorting Nexin 9), BaySNP 6743, PCR primer #2.
                                                                                                                                                                                                                                                                                             Sequence 17 BP; 1 A; 8 C; 4 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 1; Page 69; 182pp; English.
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ADF13468 standard; DNA; 17 BP.
                                                                                                                                                                                                                                                                                                                                                                                                               117 AGCGGGGGGAAAAGCC 132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14-FEB-2003; 2003WO-EP001514
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     16 AGCGGGCGGTAGAGCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                         14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          infarction and stroke.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Schwere S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-712738/67.
                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (FARB ) BAYER AG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
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Best Local S
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The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an entitinflammatory servoid and ubiquinone. A composition of the invention has antiinflammatory storaid and upiquinone, a composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, communosuppressive, and cytostatic activity. The composition may have a cuse in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of or reducing benefitivity to adenosine, reducing bronchodilation, increasing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at fib. wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human, antisense, lung dysfunction, nasal airway dysfunction, antiinflammatory steroid, ubiquinone, antiinflammatory; antiallergic; antiasthamic; hypotensive; cytostartc; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
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                                                       Query Match 2.8%; Score 12.8; DB 1; Length 17; Best Local Similarity 87.5%; Pred. No. 4e+02; Matches 14; Conservative 0; Mismatches 2; Indels
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                   Seguence 17 BP; 1 A; 11 C; 2 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; SEQ ID NO 11632; 872pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           lung inflammation; respiratory disease; ds.
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L, Shahabuddin S;
                                                                                                                                                                                                                                                                                                                                                                                                                           Human C/EBP antisense fragment no.2250.
                                                                                                                                               334 GGGGCGAGGCGAGGT 349
                                                                                                                                                                                                                                                                                               BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (EPIG-) EPIGENESIS PHARM INC
                                                                                                                                                                                                                                                                                               ABZ96390 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                          GGGCGAGGCCAGGGT
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Tang L,
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Miller S,
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The invention comprises nucleic acids (e.g. antisense oligonucleotides)

that down regulate the expression or inhibit the function of a receptor

for a neurite growth inhibitor, NOGO, prostaglandin D2 receptor (PTGDR),

KappaB kinase (IKK), or protein kinase PKR. The nucleic acids of the

invention are useful for treating; cerebrovascular accident, central

nervous system (CKS) injury, spinal cord injury, cancer (e.g. melanoma,

Imphoma or glioma), inflammatory disease (e.g. rhemmatoid arthritis,

restenosis or asthma), Crohn's disease, diabetes, obesity, autoimmune

disease, lupus, multiple sclerosis, transplant/graft rejection,

ischemeia/reperfusion injury, glomerulonephritis, sepsis, and allergic

conditions (e.g. asthma, allergic rhinitis or atopic dermatitis). The

nucleic acids of the invention are also useful for down-regulating the

cypression of a target gene and as a diagnostic tool to examine genetic

drifts and mutations within diseased cells or to detect the presence of a

target RNA in a cell. The present RNA sequence represents a human IKK-

gamma substrate sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         antisense oligonucleotide; neurite growth inhibitor; NOGO; prostaglandin D2 receptor; PTGDR; IkappaB kinase; IKK; protein kinase PKR, cerebrovascular accident; central nervous system injury; CNS injury; spinal cord injury; cancer; melanoma; lymphoma; glioma; inflammatory disease; rheumatoid arthritis; restenosis; asthma; Crohn's disease; diabetes; obesity; autoimmune disease; lupus; multiple sclerosis; transplant rejection; graft rejection; ischaemia; reperfusion; glomerulonephritis; sepsis; allergic rhinitis; atopic dermatitis; Human IKK-gamma;
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                                                                                                                                                                                               Gaps
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                                                                                                              DB 1; Length 17;
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                                  Sequence 17 BP; 0 A; 12 C; 5 G; 0 T; 0 U; 0 Other;
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                                                                                                       ch 2.8%; Score 12.8; DB 1.1 Similarity 87.5%; Pred. No. 4e+02; 14; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human IKK-gamma substrate sequence #1197.
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                                                                                                                                                                                                                                                    257 GCGGTCGGCCCGGGGC 272
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADL48687 standard; RNA; 17 BP.
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29-MAY-2001; 2001US-0294412P.
28-AUG-2001; 2001US-0315315P.
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                                                                                                                                             Best Local Similarity
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The invention comprises nucleic acids (e.g. antisense oligonucleotides)

that down regulate the expression or inhibit the function of a receptor

for a neurite growth inhibitor, NOGO, prostaglandin Dz receptor (PTGDR),

KappaB kinase (IKK), or protein kinase PKR. The nucleic acids of the

invention are useful for treating: cerebrovascular accident, central

nervous system (CNS) injury, spinal cord injury, cancer (e.g. melanoma,

lymphoma or glioma), inflammatory disease (e.g. rheumatoid arthritis,

restenosis or asthma), Crohn's disease, diabetes, obesity, autoimmune

ischaemia/reperfusion injury, glomerulonephritis, sepsis, and allergic

conditions (e.g. asthma, allergic rhinitis or atopic dermatitis). The

nucleic acids of the invention are also useful for down-regulating the

expression of a target gene and as a diagnostic tool to examine genetic

drifts and mutations within diseased cells or to detect the presence of a

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                                                                                                                                                                                                                                                                                                                                                                                                         antisense oligonucleotide; neurite growth inhibitor; NOGO; prostaglandin D2 receptor; PTGDR; IkappaB kinase; IKK; protein kinase PKR; cerebrovascular accident; central nervous system injury; CNS injury; spinal cord injury; cancer; melanoma; lymphoma; glioma; inflammatory disease; rheumatoid arthritis; restemosis; astima; Crohn's disease; diabetes; obesity; autoimmune disease; lupus; multiple sclerosis; transplant rejection; graft rejection; ischaemia; reperfusion; glomerulonephritis; sepsis; allergy; asthma; allergic rhinitis; atopic dermatitis; human PTGDR;
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                                 2.8%; Score 12.8; DB 1; Length 17; 75.0%; Pred. No. 4e+02; Live 2; Mismatches 2; Indels
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Sequence 17 BP; 2 A; 5 C; 8 G; 0 T; 2 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 161; SEQ ID NO 5095; 317pp; English.
                                                                                                                                                                                                                                                                                                                                                                        Human PTGDR substrate sequence #681.
                                                                                                                246 GCCTGGAGGCCGCGT 261
                                                                                                                                                                                                                                                     ADL51562 standard; RNA; 17 BP.
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29-MAY-2001; 2001US-0294412P.
28-AUG-2001; 2001US-0315315P.
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                                                                        12; Conservative
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that down regulate the expression or inhibit the function of a receptor
for a neurite growth inhibitor, NOGO, prostaglandin DZ receptor (PTGDR),
Ikappa kinase (IKK), or protein kinase PKR. The nucleic acids of the
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nervous system (CKS) injury, spinal cord injury, cancer (e.g. melanoma,
Ikappa or glioma), inflammatory disease (e.g. rheumatoid arthritis,
restenosis or asthma), Crohn's disease, diabetes, obesity, autoimmune
disease, lupus, multiple sclerosis, transplant/graft rejection,
is schemnia/reperfusion injury, glomearilonephritis, sepsis, and allergic
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substrate sequence.
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                           DB 1; Length 17;
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                                                      2; Indels
Sequence 17 BP; 0 A; 6 C; 8 G; 0 T; 3 U; 0 Other;
                          2.8%; Score 12.8; DB 1
8.8%; Pred. No. 4e+02;
ve 3; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 161; SEQ ID NO 5060; 317pp; English
                                                                                                                                                                                                                                                               Human PTGDR substrate sequence #646.
                                                                                                                                                                              ADL51527 standard; RNA; 17 BP
                                                                                182 GCTGCTGGCCCGTTCG 197
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29-MAY-2001; 2001US-0294412P.
28-AUG-2001; 2001US-0315315P.
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                                        68.88;
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                                        Best Local Similarity 68.8
Matches 11; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                      substrate; ds
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that down regulate the expression or inhibit the function of a receptor that down regulate the expression or inhibit the function of a receptor (PTGDR).

IkappaB kinase (IKK), or protein kinase PRR. The nucleic acids of the invention are useful for treating: cerebrovascular accident, central nervous system (CNS) injury, spinal cord injury, cancer (e.g. melanoma, lymphoma or glioma), inflammatory disease (e.g. rheumatoid arthritis, restenosis or asthma), Crohn's disease, diabetes, obesity, autoimmune disease, lupus, multiple sclerosis, transplant/graft rejection, ischaemia/reperfusion injury, glomerulonephritis, sepsis, and allergic conditions (e.g. asthma, allergic rhintis or atopic dermattitis). The nucleic acids of the invention are also useful for down-regulating the expression of a target gene and as a diagnostic tool to examine general carget RNA in a cell. The present RNA sequence represents a human NOGO receptor amberzyme substrate sequence.
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                                                                      2.8%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 4e+02;
                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human NOGO receptor amberzyme substrate sequence #59.
G; 0 T; 0 U; 0 Other;
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                                                                                                                                             0; Mismatches
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                                                                                                                                                                                                                389 CCCCGCGCGCGCGCG 404
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADM09504 standard; RNA; 17 BP.
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29-MAY-2001; 2001US-0294412P.
28-AUG-2001; 2001US-0315315P.
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   Sequence 17 BP; 2 A; 7 C;
                                                                                                                                                                                                                                                                                      CTCCGCGCGCGGCTCG
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                                        Query Match
Best Local Similarity 87.5.
"..rches 14; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2003-058513/05.
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The invention comprises nucleic acids (e.g. antisense oligonucleotides)
that down regulate the expression or inhibit the function of a receptor
for a neurite growth inhibitor, NOGO, prostaglandin D2 receptor (FYGDR),
IkappaB kinase (IKK), or protein kinase PKR. The nucleic acids of the
invention are useful for treating: cerebrovascular accident, central
nervous system (CNS) injury, spinal cord injury, cancer (e.g. melanoma,
Iymphoma or glioma), inflammatory disease (e.g. rheumatorid arthritis,
restenosis or asthma), Crohn's disease, diabetes, obesity, autoimmune
disease, lupus, multiple sclerosis, transplant/graft rejection,
ischammia/reperfusion injury, glomerulonephritis, sepsis, and allergic
conditions (e.g. asthma, allergic rhinitis or atopic dermatitis). The
nucleic acids of the invention are also useful for down-regulating the
expression of a target gene and as a diagnostic tool to examine genetic
drifts and mutations within diseased cells or to detect the presence of a
target RNA in a cell. The present RNA sequence represents a human IKK-
gamma substrate sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                               antisense oligonucleotide; neurite growth inhibitor; NOGO; prostaglandin D2 receptor; PTGDR; IkappaB kinase; IKK; protein kinase PKR; cerebrovascular accident; central nervous system injury; CNS injury; spinal cord injury; cancer; melanoma; lymphoma; glioma; inflammatory disease; rheumatoid arthritis; restennsis; asthma; Crohn's disease; diabetes; obesity; autoimmune disease; lupus; multiple sclerosis; transplant rejection; graft rejection; ischaemia; reperfusion; glomerulonephritis; sepais; allergy; asthma; allergic rhinitis; atopic dermatitis; Human IKK-gamma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel enzymatic nucleic acid that down-regulates expression of neurite growth inhibitor receptor, prostaglandin D2 receptor, IkappaB kinase or protein kinase PKR genes, for treating cancer and inflammatory disease.
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                                           DB 1; Length 17;
                                                                                 2; Indels
Sequence 17 BP; 0 A; 6 C; 8 G; 0 T; 3 U; 0 Other;
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                                      2.8%; Score 12.8; DB 1
75.0%; Pred. No. 4e+02;
tive 2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                            Human IKK-gamma substrate sequence #376.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 59; SEQ ID NO 1399; 317pp; English
                                                                                                                          263 GGCCCGGGGCTTCTCC 278
                                                                                                                                                                                                                                                                      ADL47866 standard; RNA; 17 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              05-APR-2001; 2001US-00827395.
29-MAY-2001; 2001US-0294412P.
28-AUG-2001; 2001US-0315315P.
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                                                                                 12; Conservative
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                                                              Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          substrate; ds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unidentified
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                                      Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             nucleic acid molecule in a manner that allows its expression), a mammalian cell including the expression vector and an enzymatic nucleic acid molecule that cleaves RNA derived from a GRID gene. The nucleic acid molecule is useful for treating a condition associated with the level of GRID, e.g. tissue/graft rejection and leukaemia. The present sequence is a target region for the enzymatic nucleic acids of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New nucleic acid molecule that down-regulates expression of Grb2-related with insert domain (GRID) gene, useful for treating a condition associated with the level of GRID, e.g. tissue/graft rejection and
                                                                                                                                                                                                                                                                                                                                                                                                                    Human, 88, GRID, Grb2-related with insert domain, hammerhead ribozyme,
NCH ribozyme, G-cleaver ribozyme, Zinzyme, DNAzyme, amberzyme, Inozyme,
hairpin ribozyme, tissue rejection, graft rejection, leukaemia.
                                                                                               Gaps
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                                                     DB 1; Length 17;
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                 Sequence 17 BP; 1 A; 6 C; 8 G; 0 T; 2 U; 0 Other;
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                                                                       4e+02;
                                                   Query Match 2.8%; Score 12.8; E
Best Local Similarity 75.0%; Pred. No. 4e+0
Matches 12; Conservative 2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                Human GRID mRNA substrate sequence #331.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 4; SEQ ID NO 331; 74pp; English.
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                                                                                                                                                                                                                                                                   ADM54056 standard; mRNA; 17
                                                                                                                                 249 TGGAGGCCGCGGTCGG
                                                                                                                                                                                                                                                                                                                                            (first entry)
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MCSWIGGEN J.
HAMBLIN P A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2003-829646/77.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       JARV/) JARVIS T.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
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(MCSW/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This invention describes a novel composition (a) a first active agent, comprising oligonuclectides, effective for alleviating bronchoconstruction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonuclectides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung alrway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonuclectides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallargic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated
                            ö
                                                                                                                                                                                                                                                                          respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antialergic; antiinflammatory; antialantacts; analysesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory diserses syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ds.
                            Gaps
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                                                                                                                                                                                                                                                               antisense; bronchoconstriction; allergy; hyposecretion; pain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted tnucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pabalan J, Aguilar D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       with a disease or condition such as pulmonary vasoconstriction,
                            ö
2.8%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 4e+02; ative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 15; SEQ ID NO 11632; 763pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Katz E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Li Y, Sandrasagra A, Ka
Tang L, Shahabuddin S;
                                                                                                                                                                                                                                  Human C/EBPN DNA fragment 2250.
                                                                                                                                                   ВР
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (EPIG-) EPIGENESIS PHARM INC
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ABD20299 standard; DNA; 17
                                                       CCTCCCGGGGACCTGC
                                                                                                                                                                                                       (first entry)
                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   bronchodilating agent
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              Similarity
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                                                                                                                                                                                                        29-JUL-2004
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Query Match
Best Local (
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              Best Loc
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inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary hypertension, emplysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Diagnosing glioma by detecting expression product of any one of 255 genes, glioma endothelial markers, in brain tissue sample suspected of being neoplastic, and comparing the expression with expression in normal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gape
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             marker; GEM;
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                                                                                                                                                                                                                                                                                                                                                                                                                           2.8%; Score 12.8; DB 1; Length 17;
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0; Mismatches
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01-APR-2003; 2003US-0458978P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
nes 14; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO2004016758-A2.
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Disclosure, SEQ ID NO 1053; 122pp; English.

The invention relates to an enzymatic nucleic acid molecule that percifically cleaves RNA derived from hepatitis B virus (HBV) and comprising one or more binding arms, without requiring the presence of a 2.-OH group within the molecule for activity. The nucleic acids are useful for treating hepatitis B virus infection, hepatitis, either alone or in combination with other theraples such as lamivudine and interferons. The mucleic acids are useful as diagnostic tools to examine genetic drift and mutations within diseased cells, for detecting the presence of HBV RNA in a cell, for the study of RNA and for down-regulating gene expression of a cell, for the study of RNA and for down-regulating gene expression of sequence represents an HBV RNA target sequence, used in the scope of the invention. Note: The sequence data for this patent is also available in electronic format from USPTO at sequence.

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Gaps

.; 0

2.8%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 4e+02; tive 0; Mismatches 2; Indels

117 AGCGGGGGGAAAAGCC 132

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Local Similarity 87.5

Matches

Query Match

Sequence 17 BP; 1 A; 8 C; 4 G; 0 T; 4 U; 0 Other;

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dentifying test compound as a potential anticancer drug if it decreases the expression of at least one gene; (3) identifying (M4) a test compound as a potential anticancer or antiglioma drug involves contacting a test compound at particular anticancer or antiglioma drug involves contacting a test compound with the cell which expresses mRNA of at least one gene identifying the test compound as a potential anticancer drug if it decreases the expression of at least one gene; and (4) inducing MNS of the gene, and identifying the test compound as a potential anticancer drug if it decreases the expression of at least one gene; and (4) inducing (M5) and immune response to glioma involves administering to a mammal, a protein or (1). (1) have cytostatic activities, and can be used to trigger immune destruction of glioma cells, and as immune response inducers. (M1) is useful for aiding in diagnosing glioma. (M2) is useful for treating multiple of dipma in a human. (M5) is useful for inducing an immune response to a glioma in a human. (M5) is useful for inducing an immune response to a glioma in a mammal having glioma or in a mammal who has had a glioma surgically removed. The present sequence represents a human GEM contact.
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                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hepatitis B virus, HBV; ss, enzymatic nucleic acid; RNA cleavage; hepatitis B virus infection; hepatitis; hepatocellular carcinoma; cirrhosis; liver failure; lamivudine; interferon; genetic drift; virucide; hepatotropic; antiinflammatory; cytostatic.
                                                                                                                                                                                                                                                                                                                                                                                                                              ;
                                                                                                                                                                                                                                                                                                                                                                                  2.8%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 4e+02; 1ive 0; Mismatches 2; Indel8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hepatitis B virus (HBV) RNA target sequence #1053.
                                                                                                                                                                                                                                                                                                                                               Sequence 17 BP; 1 A; 8 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14-MAY-1992; 92US-00882712.

07-FEB-1994; 94US-00193627.

08-NOV-1999; 90US-00436430.

20-MAR-2000; 2000US-00531025.

08-JUN-2000; 2000US-00636385.

24-OCT-2000; 2000US-00696347.

08-JUN-2001; 2001US-00696347.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           177 TGTCAGCTGCTGGCCC 192
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADM58919 standard; RNA; 17 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity 87.5
les 14; Conservative
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Enzymatic nucleic acid molecules which specifically cleave RNA derived from hepatitis C virus (HCV), useful for the treatment of HCV infections,
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                                                                                                                                                                                               ss; enzymatic nucleic acid; RNA cleavage; hepatitis C virus; HCV; HCV infection; type I interferon; DNAzyme.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Macejack D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 from hepatitis C virus (HCV), useful for the treatment of especially in combination with type I interferon therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pavco PA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; SEQ ID NO 917; 198pp; English.
                                                                                                                                                                  HCV DNAzyme substrate sequence #917.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Roberts E,
                                                                               ADI83671 standard; RNA; 17 BP
                                                                                                                                                                                                                                                                                                                            18-DEC-2000; 2000US-00740332.
                                                                                                                                                                                                                                                                                                                                                         18-DEC-2000; 2000US-00740332.
                                                                                                                                       (first entry)
16 AGCGGCGGTAGAGCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mcswiggen J,
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MCSWIGGEN J.
ROBERTS E.
PAVCO P A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                              MACEJACK D.
                                                                                                                                                                                                                                          Hepatitis C virus.
                                                                                                                                                                                                                                                                     US2003125270-A1.
                                                                                                                                       03-JUN-2004
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                                                                                                            ADI83671;
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(PAVC/)
(MACE/)
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                                                   RESULT 473
                                                                 ADI83671
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Novel enzymatic nucleic acid molecule such as DNAzymes and inozymes specifically cleaving RNA derived from hepatitis B virus and comprising one or more binding arms, useful for treating hepatitis and cirrhosis.

Morrissey D;

Mcswiggen JA,

Blatt L,

Draper K,

WPI; 2004-247781/23.

BLATT L. MCSWIGGEN J A. MORRISSEY D.

(BLAT/) (MCSW/) (MORR/)

DRAPER K.

(DRAP/)

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enzymatic nucleic acid; RNA cleavage; hepatítis C virus; HCV; infection; type I interferon; DNAzyme.
                                                                                       HCV DNAzyme substrate sequence #1425
                                                                                                                                                                                                                                    18-DEC-2000; 2000US-00740332.
                                                                                                                                                                                                             18-DEC-2000; 2000US-00740332.
                       ADI84179 standard; RNA; 17
                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                            Mcswiggen J,
                                                                                                                                                                                                                                                                   (MCSW/) MCSWIGGEN J.
(ROBE/) ROBERTS E.
(PAVC/) PAVCO P A.
(MACE/) MACEJACK D.
                                                                                                                                                                                                                                                                                                                                                 WPI; 2004-031273/03.
                                                                                                                                             Hepatitis C virus
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         ADI84179
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                                                                             Gaps
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interferons. The present sequence represents a HCV DNAzyme substrate
                                                                                                                                                                                                                                                                    enzymatic nucleic acid; RNA cleavage; hepatitis C virus; HCV; infection; type I interferon; DNAzyme.
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                                                    2.8%; Score 12.8; DB 1; Length 17; 75.0%; Pred. No. 4e+02; tive 2; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Macejack D;
                                Sequence 17 BP; 1 A; 9 C; 5 G; 0 T; 2 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pavco PA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; SEQ ID NO 3639; 198pp; English.
                                                                                                                                                                                                                                               HCV DNAzyme substrate sequence #3639.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Roberts E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        223
                                                                                                210 ACCTGCGGCGGGTCGC 225
                                                                                                                                                                             ADI86393 Btandard; RNA; 17 BP
                                                                                                                ACCUGCGGCCGCUCGC 17
                                                                                                                                                                                                                                                                                                                                                                        18-DEC-2000; 2000US-00740332
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                                                                                                                                                                                                                         (first entry)
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                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mcswiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                  BLATT L.
MCSWIGGEN J.
ROBERTS E.
PAVCO P A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2004-031273/03.
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nes 14; Conserv
                                                                 Similarity
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                                                                                                                                                                                                                                                                                                                             JS2003125270-A1.
                                                                             12;
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(ROBE/) R
(PAVC/) R
(MACE/) R
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Matches
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Macejack

Pavco PA,

Roberts E,

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from hepatitis C virus (HCV), useful for the treatment of HCV infections, especially in combination with type I interferon therapy.
                                                                                                                                          The invention relates to an enzymatic nucleic acid molecule which specifically cleaves RNA derived from hepatitis C virus (HCV), in which the binding arms of the enzymatic nucleic acid molecule comprises sequences complementary to any of the defined substrate sequences given in the specification. The nucleic acid molecule may be administered for the treatment of HCV infections, especially in combination with type I interferons. The present sequence represents a HCV DNAzyme substrate
Enzymatic nucleic acid molecules which specifically cleave RNA derived
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gapa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ss; enzymatic nucleic acid; RNA cleavage; hepatitis C virus; HCV;
HCV infection; type I interferon; DNAzyme.
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                                                                                                                                                                                                                                                                                                                                                                                Sequence 17 BP; 1 A; 1 C; 11 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2; Mismatches
                                                                                                   Claim 1; SEQ ID NO 1425; 198pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HCV DNAzyme substrate sequence #3755.
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Best Local Similarity
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US09436060A-16.rng.sl

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WPI; 2004-340941/31
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es 14; Conserv
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Synthetic.
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                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 478
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                                                                                                                                                                                                        Enzymatic nucleic acid molecules which specifically cleave RNA derived from hepatitis C virus (HCV), useful for the treatment of HCV infections, especially in combination with type I interferon therapy.
                                                                                                                                                                                                                                                                        The invention relates to an enzymatic nucleic acid molecule which specifically cleaves RNA derived from hepatitis C virus (HCV), in which the binding arms of the enzymatic nucleic acid molecule comprises sequences complementary to any of the defined substrate sequences given in the specification. The nucleic acid molecule may be administered for the treatment of HCV infections, especially in combination with type I interferons. The present sequence represents a HCV DNAzyme substrate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mutant cell identification-related mutagenic oligonucleotide SeqID752.
                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cell identification; oligonucleotide-directed sequence alteration; selectable phenotype; transgenic plant; herbicide resistance; sterile plant; abiotic stress tolerance; albino plant; amino acid production; ss.
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                                                                                                                                                                                                                                                                                                                                                                                              / Match 2.8%; Score 12.8; DB 1; Length 17; Local Similarity 87.5%; Pred. No. 4e+02; les 14; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                    Macejack D;
                                                                                                                                                                                                                                                                                                                                                                           Sequence 17 BP; 1 A; 6 C; 6 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                  Pavco PA,
                                                                                                                                                                                                                                                   Claim 1; SEQ ID NO 3755; 198pp; English.
                                                                                                                                                                  Roberts E,
                                                                                                                                                                                                                                                                                                                                                                                                                                         433 GGACTCGGCTCACACA 448
                                                    .8-DEC-2000; 2000US-00740332
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADN44083 standard; DNA; 17 BP
                                                                         18-DEC-2000; 2000US-00740332
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                                                                                                                                                                  Mcswiggen J,
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                                                                                               BLATT L.
MCSWIGGEN J.
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                                                                                                                    ROBERTS E.
PAVCO P A.
                                                                                                                                           MACEJACK D
          US2003125270-A1.
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(ROBE/) 1
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(MACE/) 1
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This invention relates to a novel method of identifying a cell having a desired oligonucleotide-directed sequence alteration at a first nucleic acid target site within the cell. The method comprises identifying the desired sequence alteration in cells that have been selected for the presence of a selectable phenotype conferred by a concurrent oligonucleotide-directed sequence alteration at a second nucleic acid target site within the cells. The method is useful in identifying a cell having a desired oligonucleotide-directed sequence alteration at a first the production of plants with herbicide resistance, male or female sterile plants, abiotic stress tolerance, albino plants or plants with an acid production as well as for use in mammalian cell lines.
                                                                                                                                                                                                             Identifying a cell with a desired oligonucleotide-directed sequence alteration at a nucleic acid target site within the cell by identifying the desired sequence alteration in cells selected for the presence of a selectable phenotype.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present sequence is that of a mutagenic oligonucleotide which was used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mutant cell identification-related mutagenic oligonucleotide SeqID751.
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0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                             Example 24; SEQ ID NO 752; 303pp; English.
(NAPR-) NAPRO BIO THERAPEUTICS INC.
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(NAPR-) NAPRO BIO THERAPEUTICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               425
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                                                                       Kmiec EB, Van Brabant
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                                                                                                                             This invention relates to a novel method of identifying a cell having a desired oligonucleotide-directed sequence alteration at a first nucleic acid target site within the cell. The method comprises identifying the desired sequence alteration in cells that have been selected for the presence of a selectable phenotype conferred by a concurrent oligonucleotide-directed sequence alteration at a second nucleic acid target site within the cells. The method is useful in identifying a cell having a desired oligonucleotide-directed sequence alteration at a first nucleic acid target site within the cell. The method may be useful for the production of plants with herbicide resistance, male or female sterile plants, abiotic stress tolerance, albino plants or plants with altered amino acid production as well as for use in mammalian cell lines. The present sequence is that of a mutagenic oligonucleotide which was
Identifying a cell with a desired oligonucleotide-directed sequence alteration at a nucleic acid target site within the cell by identifying the desired sequence alteration in cells selected for the presence of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Selecting an animal for desired genotypic or potential phenotypic properties such as muscle mass and/or fat deposition, comprises testing for a single nucleotide polymorphism in intron 3 of the IGF2 gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Anorectic; Antidiabetic; Muscular; Gene Therapy; CpG island; IGF2 gene intron 3; muscle mass; fat deposition; teat number; obesity; muscle deficiency; diabetes; SNP; single nucleotide polymorphism; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Porcine TSSC5 intron 1 DNA sequence polymorphism oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /standard_name= "Single_nucleotide_polymorphism"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            2.8%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 4e+02; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 17 BP; 2 A; 5 C; 7 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                The present sequence is that of a mutagenic oused in the exemplification of the invention.
                                                                                           Example 24; SEQ ID NO 751; 303pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADO80740 standard; DNA; 17 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         410 CTGAGCTGTGGGACGT 425
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity 87.5
Matches 14; Conservative
                                                           selectable phenotype
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(GENT-) GENTEC BV.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
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The present invention relates to a method (M1) for selecting an animal for having desired genotypic or potential phenotypic properties. (M1) comprises testing the animal for the presence of a nucleic acid modification affecting: the activity of an evolutionary conserved CpG island located in intron 3 of an IGF2 gene; and/or binding of a nuclear factor to an IGF2 gene. The nuclear factor is capable of binding to a stretch of nucleotides which in the wild type pig, mouse or human IGF2 gene is part of an evolutionary conserved CpG island, located in intron 3 of the IGF2 gene. The stretch is functionally equivalent to (ADQ80709). The nucleic acid modification in ADQ80709 comprises a G to A translition at IGF2-intron3-nt3072. (M1) is useful for selecting an animal with properties related to muscle mass, fat deposition, and/or test number. CC Also claimed is a method (M2) for modulating mRNA transcription of an IGF2 gene and an embod of an IGF2 gene and/or modulating binding of a nuclear factor to an IGF2 gene. Also claimed is a method (M3) for identifying a compound capable of modulating mRNA transcription of an IGF2 gene and/or modulating binding of a nuclear factor to an IGF2 gene and/or modulating binding of a nuclear factor to an IGF2 gene and/or modulating binding of a nuclear factor to an IGF2 gene and/or modulating binding of a nuclear factor to an IGF2 gene and/or modulating binding of a nuclear factor to an IGF2 gene and/or modulating binding of a nuclear factor to an IGF2 gene and/or modulating binding of a nuclear factor to an IGF2 gene and/or modulating binding of a nuclear factor to an IGF2 gene and/or modulating binding of a nuclear factor to an IGF2 gene and/or modulating a porcine sequence tagged sites (STS) compromes capable of modulating a porcine sequence tagged sites (STS) comprising a boreity. Muscle deficiencies and diabetes. The present sequence is a porcine sequence tagged sites (STS) compressing an engagence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    small interfering RNA; siRNA;
androgen signal transduction pathway protein;
androgen signal transduction; androgen receptor; hair loss;
hyperandrogenic condition; androgenic alopecia; male pattern alopecia;
acne vulgaris; seborrhoea; female hirsutism; prostatic hypertrophy; ds.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2.8%; Score 12.8; DB 1; Length 17;
87.5%; Pred. No. 4e+02;
tive 0; Mismatches 2; Indels
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/note= "2 deoxynucleotide overhang"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 17 BP; 0 A; 14 C; 2 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Androgen receptor siRNA sense strand, SEQ ID 338.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
                      3xample 1, Page 20; 38pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADQ92762 standard; RNA; 17 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GGGGGCCGGGGAGGGG
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Best Local Similarity 87.5
Matches 14; Conservative
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                                                                                                                                                                                                                 The present invention relates to novel small interfering RNAs (siRNAs), comprising a double stranded RNA, where one strand comprises a partial nucleic acid sequence of an androgen signal transduction pathway protein, and where the double-stranded RNA inhibits translation of mRNA encoding the nucleic acid sequence of the androgen signal transduction pathway protein thereby blocking the androgen signal transduction pathway. The androgen signal transduction pathway protein is chosen from isozymes I androgen signal transduction pathway protein is chosen from isozymes I creceptor (ADQ92571), aromatase (ADQ92856), 3-beta-hydroxysteroiddehydrogenase (ADQ932896), 3-beta-hydroxysteroiddehydrogenase (ADQ93182), and steroid sulfatase hydroxysteroiddehydrogenase (ADQ93182), and steroid sulfatase hydroxysteroiddehydrogenase (ADQ93720), and steroid sulfatase hydroxysteroiddehydrogenase (ADQ93722), and steroid sulfatase (ADQ93770). The siRNAs of the invention are useful for reducing hair loss in a mammal which involves contacting several mammal shair calls with the translation of mRNA of the siRNA, where the siRNA interferes with the translation of mRNA of the androgen signal transduction protein. The siRNAs are useful for treating and peaced and account of the invention are useful for treating and opecia, acme vulgaris, seboratose, and female hirsutism and prostatic and prostatic of hyperatrophy. The present sequence is the sense strand for one such siRNA of the invartion of the invartion of the siRNA of the invartion of the siRNA of the invartion of the siRNA of the invartion and female hirsutism and prostatic of hyperatrophy.
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                                                                                                  Inhibitory nucleic acid that inhibits expression of an androgen signal transduction pathway protein useful for treating hair loss, comprises a double stranded RNA having a partial sequence encoding a pathway protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Telomerase RNA; TR; promoter; cytotoxin; cancer; neoplasia; hTR; gene therapy; thymidine kinase gene; anticancer therapy; human; mutagenesis; PCR primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 17 BP; 5 A; 5 C; 4 G; 2 T; 1 U; 0 Other;
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                                                                                                                                                                                      Claim 11; Page 41; 92pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        174 AAATGTCAGCTGCTGG 189
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Best Local Similarity 87.5
Matches 14; Conservative
                                                                    WPI; 2004-561892/54,
 (GENC-) GENCIA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     the invention
                                                                                                                                                         one strand
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO9938964-A2
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                                   Kahn S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 481
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The invention relates to promoter regions from mouse and human telomerase RNA (TR) component genes. The TR gene promoter can be linked to a heterologous gene, especially a gene encoding a cyctoxin, for therapy of cancer, especially neoplasias. The telomerase is necessary for the dysregulation of the telomerase repression pathway may cause reactivation or or upregulation of the telomerase expression in cancer. Substances, identified in the methods, can be used to block transcription from the TR gene promoter through interaction of the 5' regulatory sequences. These substances, e.g. antisense oligomucleotides, transcription factors, peptide nucleic acids and factors that disrupt signal transduction, are (especially poTG2-codAupp) comprising the promoter and a viral thymidine can be used to convert a producy, e.g. gancyclovir, so that neoplasia can be controlled or treated. Direct down-regulation of telomerase RNA gene through manipulation of transcription factors may be effective anticancer therapy and the cloning of the hTR gene promoter can allows the analysis of therapeutic molecules which modulate hTR promoter and activity sequences Ask707696-521 represent PCR primers used in cloning
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
                                                                                       Mouse and human telomerase RNA gene promoters, useful for tumor specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RNA component; human telomerase; antisense oligonucleotide; infection; neuroblastoma; bladder cancer; colon cancer; prostate cancer; cantraception; sterilisation; immunosuppression; therapeutic; hTR; immune system down-regulation; anti-inflammatory therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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Pred. No. 5.3e+02;
0; Mismatches 7; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RNA component of human telomerase (hTR) antisense oligo 14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        and mutagenisis of human TR gene (hTR) promoter region
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25 cccaggcccaccrccgcaacccg 2
                                                                                                                                                                  Disclosure; Fig 12; 109pp; English.
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2.8%;
Best Local Similarity 70.8%;
Matches 17; Conservative (
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96US-00770565.
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                                           WPI; 1999-479183/40
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           08-OCT-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO9828442-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20-DEC-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20-DEC-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       02-JUL-1998
Keith WN;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAV41169;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Kim NW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 482
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Sequences shown in AAV41169 to AAV41181 represent antisense
oligonuclectides to the RNA component of human telomerase (hTR). These
antisense oligonuclectides specifically hybridise to a nucleotide
sequence within an accessible region of the hTR, but that does not
oligonuclectides may specifically be used for detection of an RNA
component of human telomerase in a sample. This is useful for diagnosing
cancer (especially neuroblastoma, bladder, colon and prostate cancer),
oligonucleotides can inhibit the telomerase activity level in a cell by
interfering with transcription of the RNA component, decreasing the half-
oligonucleotides can inhibit the telomerase activity level in a cell by
interfering with transcription of the RNA component, decreasing the half-
collent and providing progness of the telomerase activity inhibiting assembly of
the RNA component into the telomerase holoenzyme, or inhibiting telomerase of polymerase activity of telomerase of inhibiting telomerase of inhibitin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                         - used for detecting
                  New polynucleotide(s) anti:sense to human telomerase - used for detecting or inhibiting human telomerase, e.g. for treating cancers, contraception, immuno-suppression or treating infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mouse and human telomerase RNA gene promoters, useful for tumor specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Telomerase RNA; TR; promoter; cytotoxin; cancer; neoplasia; hTR;
gene therapy; thymidine kinase gene; anticancer therapy; human;
mutagenesis; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human telomerase RNA gene (hTR) promoter specific primer h112.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2.8%; Score 12.8; DB 1; Length 30; 70.8%; Pred. No. 5.4e+02; ive 0; Mismatches 7; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 30 BP; 4 A; 10 C; 8 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RES CAMPAIGN TECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                131 CCTCGGCCTGCCGCCTTCCACCGT 154
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ccrcrrccrececcreaaaceer 28
                                                                                                                         Claim 11; Page 65; 80pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      298/c
AAZ07298 standard; DNA; 38 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     99WO-GB000308
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity 70.8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1999-479183/40.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO9938964-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     29-JAN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 483
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ID AAZ0'
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gene therapy

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The invention relates to promoter regions from mouse and human telomerase

RNA (TR) component genes. The TR gene promoter can be linked to a

RNA (TR) component genes. The TR gene encoding a cytotoxin, for therapy of

cancer, especially neoplasias. The telomerase is necessary for the

currestricted proliferative capacity of many human cancers. Mutation or

currestricted proliferative capacity of many human cancers. Mutation or

currestricted proliferative capacity of many human cancers. Mutation or

currentlation of the telomerase repression pathway may cause reactivation

currentlation of the telomerase expression pathway may cause reactivation

currentlation of the telomerase expression pathway may cause reactivation

currentlation of the language or used to block transcription from the TR

currentlation for an be used to block transcription factors,

cuseful for cancer therapy. In particular, gene therapy vectors

cuseful for cancer therapy. In particular, gene therapy vectors

cuseful for cancer therapy. In particular, gene therapy vectors

cuseful for cancer therapy and the promoter and a viral thymidine

kinase gene can be used to convert a prodrug, e.g. gancyclovir, so that

cuseful for cancer therapy and the cloning of the hrw gene promoter

cuseful cancer therapy and the colning of the hrw gene promoter

cuseful cancer therapeutic molecules which modulate hrm promoter

cutivity. Sequences AAZ07696-121 represent PCR primers used in cloning

cuseful for cancer forman TR gene (hTR) promoter region
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention describes a pair of RNA molecules (R1, R2) for detecting a first subunit protein (I) of telomerase. R1 and R2 both bind to (I) and have formulae 5'-A-B-C-3' (R1) 5'-D-E-F-3' (R2) where: A and I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RNA molecules for detecting telomerase, useful for diagnosis of or metastases, can be ligated when bound to telomerase subunit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; telomerase; hTR; telomeric repeat amplification protocol; TRAP;
identification; detection; quantification; cancer; metastasis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2.8%; Score 12.8; DB 1; Length 38; 70.8%; Pred. No. 5e+02; tive 0; Mismatches 7; Indel8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 38 BP; 3 A; 7 C; 21 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Adenovirus nucleotide sequence SEQ ID NO:20.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         223 CGCCTGCCCAGCCCCCGAACCCCG 246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CCCAGGCCCACCTCCGCAACCCG 2
                 Disclosure; Fig 12; 109pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 2; Col 23; 32pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (INVI-) INVITRO DIAGNOSTICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28-JUN-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 70.8
Matches 17; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    12-JUN-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAA08205;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 484
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- RNA segment of 10 to 100000 nucleotides (nt) that together are treplicated by RNA replicase; B and B = RNA segments of 10 to 250 nt from the Y region of human telomerase and bind specifically to (1); C and D = RNA segments of 1 to 10000 nt which can be ligated together. Ligation of C and D produces R3 of formula 5'-A-B-C-D-E-F-3' (R3) with E and B bound to (1). Replication of R3 by RNA replicase indicates presence of (1). Also described are: (1) method for detecting (1) using R1 and R2; (2) kit for this process containing R1, R2, ligase and an amplification system; DNA. R1 and R2 are used to detect and quantify telomerase, particularly provide an assay that does not require expensive of makey that does not require expensive or integrity browned; and is suitable for automation. The present sequence represents an oligonucleotide used in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                  invention
8x868888888888888888888888888888
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Sequence 62 BP; 16 A; 27 C; 10 G; 9 T; 0 U; 0 Other;

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0
             Score 12.8; DB 1; Length 62;
Pred. No. 3.2e+02;
0; Mismatches 7; Indels
                                         0; Mismatches
                                                                  222 TCGCCTGCCCAGCCCCGAACCCC 245
                2.8%;
                           70.8%;
Ouery Match
Best Local Similarity 70.0.
Best Local 17; Conservative
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Gaps

61 38 rcccagcccacccrccgcaaccc AAH24816 standard; RNA; 62 485 셤

RNA-binding protein; RBP; RNA replicase; RNA identification; telomerase; Human nucleic acid sequence derived from Y-1 domain of telomerase. (first entry) Homo sapiens. 06-AUG-2001 AAH24816;

99US-00255464. 98US-0075495P. 22-FEB-1999; 29-MAY-2001

US6238867-B1

(INVI-) INVITRO DIAGNOSTICS INC.

Grossman A; Roninson IB,

WPI; 2001-366472/38.

RNA and New ribonucleic acids useful for identifying naturally occurring sequences having affinity for RNA-binding protein Example 2; Col 26; 36pp; English. components.

RNA

The specification describes a first RNA (RNA1) and a second RNA (RNA2) components. RNA1 has the formula 5'-A-B-C-3', where A is section having 10-100,000 nucleotides and is can be received by an RNA replicase and with another DNA sequence, F, being replicated; B is section having 10-3,000 nucleotides which have affinity to one consensus sequence of the RBP and which can bind to the protein component; C is section having about 1-20 nucleotides and which can be ligated to D of the second RNA molecule. RNA2 has the formula 5'-D-E-F-3', where D is section having 10 nucleotides and which can be ligated to C; E is section laving 1-20 nucleotides and which can be ligated to C; E is section 10-3,000 nucleotides which have affinity to second consensus sequence of the RBP

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The present invention describes a pair of RNA molecules (R1, R2) for detecting a first subunit protein (I) of telomerase. R1 and R2 both bind to (I) and have formulae 5'-A-B-C-3' (R1) 5'-D-E-F-3' (R2) where: A and F = RNA segment of 10 to 10000 nucleotides (nr) that together are the x region of human telomerase and bind specifically to (1); C and D = RNA segments of 1 to 10000 nt which can be ligated together. Ligation of RNA segments of 1 to 10000 nt which can be ligated together. Ligation of C and D produces R3 of formula 5'-A-B-C-D-E-F-3' (R3) with E and B bound C C and D produces R3 of formula 5'-A-B-C-D-E-F-3' (R3) with E and B bound C C and D produces R3 of formula 5'-A-B-C-D-E-F-3' (R3) with E and B bound C C and D produces R3 of formula 8'-A-B-C-D-E-F-3' (R3) with E and B bound C C and D produces R3 of formula 8'-A-B-C-D-E-F-3' (R3) with E and B bound C C and D produces R3 of formula 8'-A-B-C-D-E-F-3' (R3) with E and B bound C C and D 3' method for making R1 and R2 is ligase and an amplification system; and 3) method for making R1 and R2 by transcription from appropriate DNA. R1 and R2 are used to detect and quantify telomerase, particularly for diagnosis of cancer and for detection of metastases. R1 and R2 trained personnel, and is suitable for automation. The present sequence represents an oligonucleotide used in the exemplification of the present
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and which can bind to the protein component; F is section having 10-100,000 nucleotides which can be received by an RNA replicase and with A being replicated. RNA1 and RNA2 are capable of forming a third RNA (RNA3) of formula 5'A-B-C-D-E-F-3'. The method is useful for the identification and characterization of RNA sequences having specific affinity to amino acid consensus sequences of RBPs, and to RNAs. AAH24815-16 were used to produce a double-stranded RNA1, comprising the Y-1 domain of human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; telomerase; hTR; telomeric repeat amplification protocol; TRAP; identification; detection; quantification; cancer; metastasis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pair of RNA molecules for detecting telomerase, useful for diagnosis cancer or metastases, can be ligated when bound to telomerase subunit
                                                                                                                                                                                                                                                                                                Gaps
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0
                                                                                                                                                                                                                                           2.8%; Score 12.8; DB 1; Length 62; 70.8%; Pred. No. 3.2e+02; tive 0; Mismatches 7; Indels
                                                                                                                                                                                                  Sequence 62 BP; 16 A; 27 C; 10 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Adenovirus nucleotide sequence SEQ ID NO:19.
                                                                                                                                                                                                                                                                                                                                  222 TCGCCTGCCCAGCCCCGAACCCC 245
                                                                                                                                                                                                                                                                                                                                                                           rcccaggcccacccrccgcaccc 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 2; Col 23; 32pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                           Conservative
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                                                                                                                                                                                                                                                                Local Similarity
les 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mastadenovirus.
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                                                                                                                                                             telomerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Grossman A;
                                                                                                                                                                                                                                                                                                                                                                             38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAA08204;
                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         protein.
                                                                                                                                                                                                                                                                                    Matches
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245

222 TCGCCTGCCCAGCCCCCGAACCCC

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The specification describes a first RNA (RNA1) and a second RNA (RNA2) for use in binding an RNA-binding protein (RBP) having protein and RNA components. RNA1 has the formula 5'-A-E-C-3', where A is section having 10-100,000 nucleotides and is can be received by an RNA replicase and with another DNA sequence, F, being replicated; B is section having 10-3,000 nucleotides which have affinity to one consensus sequence of the BPP and which can bind to the protein component; C is section having about 1-20 nucleotides and which can be ligated to D of the second RNA molectides which have affinity to second consensus sequence of the RNA2 has the formula 5'-D-E-F-3', where D is section having 1-C on nucleotides which can be ligated to C; E is section 10-3,000 nucleotides which can be received by a RNA replicase and with A being replicated. RNA1 and RNA2 are capable of forming a third RNA feature S'A-B-C-D-E-F-3'. The method is useful for the identification and characterization of RNA sequences having specific affinity to amino and characterization of RNAs equences having specific affinity to amino produce a double-stranded RNA1, comprising the Y-1 domain of human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New ribonucleic acids useful for identifying naturally occurring RNA sequences having affinity for RNA-binding protein having protein and RNA
                                                                                                                                                                                                                                                                                                                                                                                    RNA-binding protein; RBP; RNA replicase; RNA identification; telomerase;
                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                  Human nucleic acid sequence derived from Y-1 domain of telomerase.
                                                                                   ;
0
                                              Score 12.8; DB 1; Length 66;
Pred. No. 3e+02;
0; Mismatches 7; Indels
                                                                               7; Indels
                  28 G; 17 T; 0 U; 0 Other;
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                                                                                                                     222 TCGCCTGCCCAGCCCCGAACCCC 245
                                                                                                                                     TCCCAGGCCCACCTCCGCAACCC 6
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                                                Query Match 2.8%;
Best Local Similarity 70.8%;
Matches 17; Conservative
                ΰ
                                                                                                                                                                                                                                            AAH24815 standard; RNA; 66
                                                                                                                                                                                                                                                                                                                 (first entry)
                  Sequence 66 BP; 10 A; 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Grossman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2001-366472/38.
                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            23-FEB-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US6238867-B1
                                                                                                                                                                                                                                                                                                                 06-AUG-2001
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The invention comprises the amino acid and coding sequences of the human cholesteryl ester transfer protein (CETP), the invention also comprises polymorphisms identified within the CETP gene. The DNA and protein sequences of the invention are useful in haplotyping and/or genotyping the CETP gene in an individual. The DNA and protein sequences may also be used to screen drugs or compounds targeting the CETP or its variant to treat a condition or disease associated with CETP (e.g. atherosclerosis, cardiovascular disease or hypercholesterolaemia). The present DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cardiovascular disease or hypercholesterolaemia). The present DNA sequence represents an allele specific oligonucleotide PCR primer for the
                                                                                                                                                                                                               human; cholesteryl ester transfer protein; CETP;
single nucleotide polymorphism; SNP; drug screening; atherosclerosis;
cardiovascular disease; hypercholesterolasemia;
allele specific oligonucleotide; ss; PCR; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New isolated polynucleotide useful for haplotyping and/or genotyping cholesteryl ester transfer protein (CETP) gene in an individual or in screening for drugs useful in treating diseases associated with CETP
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                                                                                                                                                                                  Human CETP gene allele specific oligonucleotide PCR primer #21.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Lee
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Lachowicz M,
Tirrell C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 43; SEQ ID NO 80; 250pp; English.
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Sausker EA,
   rccaggcccaccrccgcaaccc
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                                                                                                                                                                                                                                                                                                                                                                                            28-APR-2003; 2003WO-US013288.
                                                                                                                                                                                                                                                                                                                                                                                                                            26-APR-2002; 2002US-0375791P.
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                                                                                 ADG98448 standard; DNA;
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Petersen N, Rounds
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es 12; Conserv
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Gaps

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2.8%; Score 12.8; DB 1; Length 66; 70.8%; Pred. No. 3e+02; ative 0; Mismatches 7; Indels

Query Match
Best Local Similarity 70.8
Matches 17; Conservative

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Seguence:

Minimum DB Maximum DB

Searched:

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AM934303
SWOVLJCAN12H12 Onchocerca volvulus infective larva cDNA
(SAW94WL-OvL3) Onchocerca volvulus cDNA clone onch672 5' similar to TR:Q33571 Q33571 ATPASE SUBUNIT 6 ;, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Onchocerca volvulus
Onchocerca volvulus
Onchocerca volvulus
Bukaryota, Metazoas, Nematoda; Chromadorea; Spirurida; Filarioidea;
Onchocercidae; Onchocerca.

1 (bases 1 to 19)
Williams, S.A., Lizotte-Waniewski, M., Laney, S., Wenhong, L.,
Hillier, L., Allen, M., Bowles, L., Geisel, S., Jost, S., Kucaba, T.,
Martin, J., Steptoe, M., Theising, B., White, Y., Wylie, T.,
Chappell, J., Person, B., Gibbons, M., Harvey, N., Pape, D.,
Chamberlain, A., Morales, R., Schurk, R., Ritter, E., Kohn, S.,
Underwood, K. and Marta, M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ACCESSION: AA934303
ACCESSION: AA934303
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                                                                                                                                                August 24, 2005, 14:51:46; Search time 0.001 Seconds (without alignments) 17.138 Million cell updates/sec
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Smith College Department of Biological Sciences
Department of Biological Sciences, Clark Science Center, Smith
                                                                                                                                                                                                                                                                                                              1 gggttgcggagggtgggcct.....aggactcggctcacacatgc 451
GenCore version 5.1.6
(c) 1993 - 2005 Compugen Ltd.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Total number of hits satisfying chosen parameters:
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                                                                                                      - nucleic search, using sw model
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Maximum Match 100%
Listing first 2 summaries
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1 AA934303
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seg length: 200000000
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Score Match Length DB
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                               Copyright
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KEYWORDS
SOURCE
ORGANISM
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Result Š DEFINITION

RESULT 1 AA934303

ACCESSION

REFERENCE AUTHORS TITLE JOURNAL COMMENT

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Email: genome@smith.edu
The library was constructed by Wenhong Lu. The library is available from Dr. S.A. Williams, email genome@smith.edu When requesting this ollone from Dr. Williams, please reference the Williams lab clone id SWOVL3CANI2H12
- SWOVL3CANI2H12
SEQ primer: -40ml3 fwd. ET from Amersham
High quality sequence stop: 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Anotes Wedcor: lambda UniZap XR; Site 1: EcoR I; Site 2: Xho I; Cutaneous filarial nematode parasite of humans. The I; Cutaneous filarial nematode parasite of humans. The I constructed from third stage infective larvae of Onchocerca volvulus isolated from mosquitoes 10 days after infection and converted to double stranded cDNA using reverse transcriptase and oligo(dT) followed by RNase H and DNApol I. The library had 1.8 x 10E5 independent recombinants and average insert size was 900 base pairs. The library was constructed by Wenhong Lu. The library is available from Dr. S.A. Williams, email genome@smith.edu."
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SWO-KISCANI2H12 Onchocerca volvulue infective larva cDNA
(SAM94ML-OvL3) Onchocerca volvulue cDNA clone onch672 5' Bimilar to
TR:Q33571 Q33571 ATPASE SUBUNIT 6 ;, mRNA sequence.
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The library was constructed by Wenhong Lu. The library is available
from Dr. S.A. Williams, email genome@smith.edu When requesting this
clone from Dr. Williams, please reference the Williams lab clone id
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Bukaryota; Metazoa; Nematoda; Chromadorea; Spirurida; Filarioidea;
Onchocercidae; Onchocerca.

1 (bases 1 to 19)
Williams,S.A., Lizotte-Waniewski,M., Laney,S., Wenhong,L.,
Williams,S.A., Lizotte-Waniewski,M., Gaisel,S., Jost,S.,
Martin,J., Allen,M., Bowles,L., Geisel,S., Wolte,Y., Wylie,T.,
Chappell,J., Srepton,B., Gibbons,M., Harvey,N., Pape,D.,
Chamberlain,A., Worales,R., Schurk,R., Ritter,E., Kohn,S.,
Molecular Parasitology OvL3
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Contact: Steven A. Williams
Molecular Parasitclogy
Smith College Department of Biological Sciences
Department of Biological Sciences, Clark Science Center, Smith
College, Northampton, MA, 01063, USA
                                                                                                                                                                                                                                                                                                                                                                                   /mol_type="mRNA"
|strain="Sierra Leone"
|db xref="taxon|6282"
|clone="onch672"
|lab host="XL1-Blue MRF'"
|clone lib="Onchocerca volvulus infective larva cDNA
|SAM94WL-OvL33"
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    19
/organism="Onchocerca volvulus"

Northampton, MA, 01063, USA
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0; Gaps

Query Match
2.0%; Score 8.8; DB 1; Length 19;
Best Local Similarity 83.3%; Pred. No. 0;
Matches 10; Conservative 0; Mismatches 2; Indels

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	4444446664 466766			
GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compugen Ltd. ic search, using sw model gust 24, 2005, 14:28:58; Search time 1 Seconds (without alignments) 4.890 Million cell undates/sec	-436-060A-16 ttgcggagggtgggcctaggactcggctcacacat TTY_NUC 10.0 , Gapext 0.5 ags, 5421 residues satisfying chosen parameters: 504	h: 0 h: 2000000000 timum Match 0% timum Match 100% ting first 257 summaries .subdb:*	than or equal to the score of the result being print by analysis of the total score distribution. SUMMARIES y h Length DB ID Description	62 1 US-08-873-709-20 Sequence 20, Appl 10 US-08-255-464B-16 Sequence 16, Appl 20 Sequence 15, Appl 20 US-08-255-464B-15 Sequence 15, Appl 20 US-09-255-464B-15 Sequence 15, Appl 20 US-09-217-82B-1 Sequence 17, Appl 20 US-09-717-82B-1 Sequence 17, Appl 20 US-09-717-82B-1 Sequence 17, Appl 20 US-09-717-82B-1 Sequence 17, Appl 20 US-08-31 US-08-3
Copy OM nucleic - nucleic s Run on: August	Perfect score: US-09-Perfect score: 451 Sequence: 1 gggt Scoring table: IDENT) Gapop Searched: 252 ss Total number of hits	seq lengated language language Maring	score greater the and is derived keeult Query No. Score Match	C C C C C C C C C C C C C C C C C C C

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; Sequence 15, Application US/09255464B
; Patent No. 6238867
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Patent No. 6037126
GENERAL INFORMATION:
PRIOR APPLICATION NUMBER: 60/6;
PRIOR FILING DATE: 1998-02-23;
NUMBER OF SEQ ID NOS: 25
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 16
LENGTH: 62
                              1998-02-23
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Best Local Similarity 100.0
Matches 54; Conservative
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STRANDEDNESS: single
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COMPUTER READABLE FORM:
                                                                                                                                                                                                         ORGANISM: Homo sapiens
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// MOLECULE TYPE: DNA
US-08-873-709-19
                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
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                                                                                                                                                                                   TYPE: DNA
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                                 Sequence 212, App
Sequence 9, Appli
Sequence 106, App
Patent No. 5166057
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Sequence 16, Application US/09255464B

Fatent No. 6238867

GENERAL INFORMATION:

APPLICANT: Roninson, Igor

TITLE OF INVENTION: Identifying Naturally Occurring RNA Sequences Having

TITLE OF INVENTION: Affinity for RNA-Binding Proteins

FILE REPERENCE: Q001/004BE: US/09/255,464B

CURRENT FILING DATE: 1999-02-22
   Sequence 9, Appli
Sequence 212, App
Sequence 9, Appli
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OMPRATING SYSTEM: PC_DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/873,709
FLING DATE: 12-UN-1997
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Janiuk, Anthony J.
REGISTRATION NUMBER: 29,809
REGISTRATION NUMBER: 29,809
REGISTRATION NUMBER: 29,001/002
REGISTRATION NUMBER: 29,001
INFORMATION FOR EQ. 1D NO: 20:
SEQUENCE THARACTERISTICS:
LENGTH: 62 base pairs
TYPER NUCLEL CACID
TYPE NUCLEL CACID
TYPE NUCLEL CACID
TYPE NUCLE CACID
TYPE NUCLE CACID
TYPE NUCLE CACID
   US-08-871-732A-9
US-09-180-437-212
US-09-346-510B-9
US-09-544-934B-106
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Pred. No. 0.74;
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100.0%; Pred. No. v...
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ZIP: 10570
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
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COUNTRY:
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TITLE OF INVENTION: COMPOSITIONS, METHODS, KITS AND
TITLE OF INVENTION: APPARATUS FOR DETERMINING THE PRESENCE OR ABSENCE OF
TITLE OF INVENTION: PROTEIN COMPONENT OF TELOMERASE ENZYME
NUMBER OF SEQUENCES: 25
CORRESPONDENCE ADDRESS:
ADDRESSEE: Abraham Grossman
STREET: 666 Washington Avenue
CITY: Pleasantville
STATE: NY
                                                                                                               0; Сарв
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gape
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                                                                                                                                                                                                                   1 GGGTTGCGGAGGGTGGGCCTGGGAGGGGTGGTGGCCATTTTTTGTCTAACCCTA 54
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DB 1; Length 62; 0.74;
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FILING DATE: 12-JUN-1997
CLASSIFICATION NUMBER: US/08/873,709
FILING DATE: 12-JUN-1997
CLASSIFICATION NUMBER: 29,809
REFERENCE/POCKET NUMBER: 20,809
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1;
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tch 12.0%; Score 54; DB al Similarity 100.0%; Pred. No. 0.7 54; Conservative 0; Mismatches
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Gaps

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Indels

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DB 1; Length 31;
                                                                  Score 31; DB 1; Pred. No. 20; 9; Mismatches
                                                                                                                                                             40 TITIGICIAACCCIAACTGAGAAGGGCGIAG 70
                                                                                                                                                                                       1 UUUUGUCUAACCCUAACUGAGAAGGGCGUAG 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: 09-APR-1997
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                       Sequence 49, Application US/09349532 Patent No. 6294650
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 01:
TELECOMMUNICATION:
TELEPHONE: (415) 576-0200
  MOLECULE TYPE: RNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: RNA (genomic)
                                                                  Query Match 6.9%;
Best Local Similarity 71.0%;
Matches 22; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 31 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 7
US-09-717-828B-1/c
           ; MODECOME 111
US-08-838-545-49
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-09-349-532-49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
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         APPLICANT: Roninson, Igor
APPLICANT: Grossman, Abraham
ITILE OF INVENTION: Compositions, Methods, Kits and Apparatus for
ITILE OF INVENTION: Compositions, Methods, Kits and Apparatus for
ITILE OF INVENTION: Identifying Naturally Occurring RNA Sequences Having
ITILE OF INVENTION: Affinity for RNA-Binding Proteins
FILE REFERENCE: Q001/004a
CURRENT FILING DATE: 1999-02-22
PRIOR APPLICATION NUMBER: 60/075,495
PRIOR APPLICATION NUMBER: 60/075,495
PRIOR FILING DATE: 1998-02-23
NUMBER OF SEQ ID NOS: 25
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 15
LENGTH: 66
LENGTH: 66
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APPLICANT: Wright, Woodring E.
APPLICANT: Wright, Woodring E.
APPLICANT: Wright, Weodring E.
APPLICANT: Corey, David R.
APPLICANT: Corey, David R.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: 60 CORRESPONDENCE ADDRESS: 60 CORRESPONDENCE ADDRESS: ADDRESSE: Two Embarcadero Center, Eighth Floor CITY: San Francisco CITY: San Francisco CITY: California
                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 54; DB 1; Length 66;
Pred. No. 0.81;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Indels
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SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/838,545
FILING DATE: 09-APR-1997
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Stoorella, John R.
REFERENCE/DOCKET NUMBER: 015389-001610US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFROME (415) 576-0300
INFORMATION FOR SEQ ID NO: 49:
SEQUIENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 GGGTTGCGGAGGGTGGGCCTGGGAGGGGTGGTGGCCATTT
                                                                                                                                                                                                                                                                                                                                                                                                                       12.0%; Sco...
100.0%; Pred. No. v...
... 0; Mismatches
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 49, Application US/08838545
Patent No. 6046307
GENERAL INFORMATION:
APPLICANT: Shay, Jerry W.
                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 100.0
Matches 54; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 31 base pairs
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STRANDEDNESS: single
TOPOLOGY: linear
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  USA
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GENERAL INFORMATION:
APPLICANT: Shay, Jerry W.
APPLICANT: Wright, Woodring E.
APPLICANT: Platyszek, Mieczysław A.
APPLICANT: Corey, David R.
APPLICANT: Corey, David R.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 60
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cn 6.9%; Score 31; DB 1; Length 31; 1 Similarity 71.0%; Pred. No. 20; 22; Conservative 9; Mismatchen 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/09/349,532
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            015389-001610US
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NAME/KEY: misc_feature LOCATION: (31)
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CITY: Palo Alto
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OTHER INFORMATION: Biotin 5'-terminal

NAME/KEY: misc_feature

LOCATION: (31)

OTHER INFORMATION: Biotin 3'-terminal

OTHER INFORMATION: Description of Artificial Sequence: Affinity Agent
US-09-717-828B-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Wainrich, Scott L
APPLICANT: Atkinson III, Edward M
APPLICANT: Lichtsteiner; Serge P
APPLICANT: Lichtsteiner; Serge P
APPLICANT: Lichtsteiner; Serge P
APPLICANT: Usescrit, Allain P
APPLICANT: Pruzan, Ronald A
TITLE OF INVENTION A Method for Purifying Telomerase
CURRENT APPLICATION NUMBER: US/09/11/B29A
CURRENT FILING DATE: 1999-10-18
PRIOR FILING DATE: 1999-10-18
PRIOR FILING DATE: 1999-10-18
PRIOR PELICATION NUMBER: 08/813,377
PRIOR PELING DATE: 1995-08-04
NUMBER OF SEQ ID NOS: 11
SOFTWARE: PatentIn Ver: 2.1 edited
SEQ ID NO 1
LENGTH::31
                                                           APPLICANT: Wathrich, Scott L
APPLICANT: Atkinson III, Edward M
APPLICANT: Lichteiner, Serge
APPLICANT: Lichteiner, Serge
APPLICANT: Lichteiner, Serge
APPLICANT: Lichteiner, Serge
APPLICANT: Pruzan, Ronald A
TITLE OF INVENTION: A Method for Purifying Telomerase
CURRENT APPLICATION NUMBER: US/09/717,828B
CURRENT APPLICATION NUMBER: 09/420,056
PRIOR FILING DATE: 1999-10-18
PRIOR FILING DATE: 1999-10-18
PRIOR FILING DATE: 1999-04-04
PRIOR FILING DATE: 1995-08-04
PRIOR FILING DATE: 1995-08-04
PRIOR FILING DATE: 1995-08-04
INUMBER OF SEQ ID NOS: 11
SOFTWARE: Patentin Ver. 2.1 edited
IBNGTH: 31
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       31 Trercraacceraacreagaagagagagaga 1
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LOCATION: (1)
OTHER INFORMATION: Biotin 5'-terminal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 1, Application US/09717829A Patent No. 6545133
Sequence 1, Application US/09717828B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: DNA
ORGANISM: Artificial Seguence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 31; Conservative
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US-09-717-829A-1/C
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APPLICANT: Weinrick, Scott
APPLICANT: Weinrich, Scott
APPLICANT: Weinrich, Scott
APPLICANT: Weinrich, Scott
APPLICANT: Weinson III, Edward
APPLICANT: Alain
APPLICANT: Vasserot, Alain
APPLICANT: Vasserot, Alain
APPLICANT: Vasserot, Alain
APPLICANT: Pracan, Ronald
TITLE OF INVENTION: Unhibitors
TITLE OF INVENTION: Unhibitors
TITLE OF INVENTION: Unhibitors
TITLE OF INVENTION: Unhibitors
FILE REFERENCE: 01/06C
CURRENT APPLICATION NUMBER: 08/510,736
PRIOR PILING DATE: 1995-08-04
PRIOR PELING DATE: 1995-04-04
PRIOR APPLICATION NUMBER: 09/420,056
PRIOR PLING DATE: 1999-10-18
PRIOR PLING DATE: 2000-11-20
NUMBER OF SEC ID NOS: 11
SOFTWARE: Patentin version 3.1
FEASTION ON 1
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; OTHER INFORMATION: Biotin 3'-terminal ; OTHER INFORMATION: Description of Artificial Sequence: Affinity Agent US-09-717-829A-1
                                                                                                                                                                              Gaps
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Sequence 22, Application US/08330123A

Sequence 22, Application US/08330123A

Sequence 22, Application US/08330123A

SEQUENCE CANTAINON:

APPLICANT: VILLEPONTEAU, Bryant
APPLICANT: FUNG, Junii
APPLICANT: FUNK, Walter

APPLICANT: AUDREWS, William H.

TITLE OF INVENTION: HUMAN TELOMERASE

NUMBER OF SEQUENCES: 25

CORRESPONDENCE ADDRESS:

ADDRESSEE: TOWNSEND and Townsend Khourie and Crew

STREET: 379 Lytton Avenue
                                                                                                                                                                                                                                 42 TIGICIAACCCIAACTGAGAAGGGCGTAGGC 72
                                                                                                                  DB
20;
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                                                                                                               Query Match 6.9%; Score 31; DB Best Local Similarity 100.0%; Pred. No. 20; Matches 31; Conservative 0; Mismatches
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Patent No. 6787133
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Best Local Similarity 100.(
Matches 31; Conservative
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015389-000830US
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CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-UL-1994
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-0006
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-660-678A-22/c
; Sequence 22, Application US/08660678A
; Patent No. 5837857
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFRENCE/DOCKET NUMBER: 01536
TELECOMMUNICATION INFORMATION:
TELEPAK: (415) 576-0200
INFORMATION FOR SEQ ID NO: 22:
SEQUENCE CHARACTERISTICS:
LENGAL 30 base pairs
TENER: nucleic acid
STRANDEDNESS: single
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TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 22:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 30 base pairs
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                      linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY:
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APPLICANT: Feng, Junia
APPLICANT: Feng, Junia
APPLICANT: Funk, Walter
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Assays for the RNA Component of Human
TITLE OF INVENTION: Telomerase
TITLE OF SEQUENCES: 40
CORRESPONDENCE ADDRESS:
ADDRESSEE: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1; Length 30; 23;
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                               COMPUTER PAGABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC Compatible
COMPUTER: PatentIn Release #1.0, Version #1.25
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/330,123A
FILING DATE: 27-OCT-1994
CLASSFETACTION: 435
PRICK APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-OLL-1994
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M
REGISTRATION NUMBER: 15389-000810
TELEPHONE: (415) 326-2400
TELEFAX: (415) 326-2400
TELEFAX: (415) 326-2402
INFORMATION FOR SEQ ID NO: 22:
SEQUENCE CHARACTERISTICS:
LENGTH: 30 base pairs
TYPE: MOLAIC CHARACTERISTICS:
LYPE: MOLAI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: Patentln Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/482,115B FILING DATE: 07-JUN-1995 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  77 TGCTTTTGCTCCCCGCGCGCTGTTTTCTC 106
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APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-UUL-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-CCT-1994
ATTORNEY/AGENT INFORMATION:
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COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 11
US-08-482-115B-22/c
; Sequence 22, Application US/08482115B
; Setnent No. 5776679
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 6.7
Best Local Similarity 100.
Matches 30; Conservative
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STATE: California
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; MOLECULE TYPE:
US-08-330-123A-22
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DB 1; Length 30;
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COUNTRY: USA
COUNTRY: USA
ZIP: 94111-3834
COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/660,678A
FILING DATE: 05-UN 1996
FILING DATE: 05-UN 1996
FILING DATE: 05-UN 1996
FILING DATE: 07-UN 1996
FILING DATE: 07-UN 1996
                                                                                                                                                                                                                                                                                                                         APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junil
APPLICANT: Feng, Junil
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 30
COMMESSONDERICE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
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Methods for Detecting the RNA Component of Telomerase
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APPLICANT: Feng, Junia
APPLICANT: Feng, Junia
APPLICANT: Fung, Junia
APPLICANT: Marhuenda, Maria Antonia Blasco
APPLICANT: Willeponteau, Bryant
TITLE OF INVENTION: RNA COMPONENT OF TELOMERASE
NUMBER OF SEQUENCES: 45
CORRESPONDENCE ADDRESS:
ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C.
STREET: Two Militia Drive
                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Floppy disk
COMPUTER: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/770,565
FLING DATE: 20-DEC-1996
CLASSIFICATION: 435
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SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/485,778
FILING DATE: 07-JE-1995
                                                                                       LLP
TITLE OF INVENTION: Methods for Detecting th
TITLE OF INVENTION: Telomerase
NUMBER OF SEQUENCES: 26
NUMBER OF SEQUENCES: 26
ADDRESSEE: TOWNSEND and TOWNSEND and CREW
STREET: TOWN Embarcadero Center, 8th Floor
STREET: Can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     015389-002300US
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION UNDMBER: 32,944
REFERENCE/DOCKET NUMBER: 01536
TELECOMMUNICATION INFORMATION:
TELEFAM: 415-576-0200
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Andrews, William H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity 100 hes 30; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 30 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 02173
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET: Two man
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                                                                                                                                                                        USA
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                                                                                                                                                                      COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-770-565-8
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Matches
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APPLICANT: Wim, Nam Woo
APPLICANT: Wim, Name Woo
APPLICANT: Wealey, James T.
APPLICANT: Realey, James T.
APPLICANT: Weinrich, Scott L.
APPLICANT: Weinrich, Scott L.
APPLICANT: Weinrich Bethods for Detecting the RNA Component of TITLE OF INVENTION: Telomerase
NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS: ADDRESSE: TOWNSEND and CREW LLP
STREET: Two Embarcadero Center, 8th Floor
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    DB 1; Length 30;
23;
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23;
                                              0; Indels
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MEDIUM TYPE: Floppy disk
COMPUTER: Floppy disk
COMPUTER: Ploppy disk
COMPUTER: PLOPPY disk
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/770,565
FILING DATE: 20-DEC-1996
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE DOCKET NUMBER: 015389-002300US
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-576-0200
TELEFAX: 415-576-0300
                                                                                       TTCTC 106
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  6.7%; Score 30; DB 100.0%; Pred. No. 23; tive 0; Mismatches
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Patent No. 5846723
GENERAL INFORMATION:
APPLICANT: Wu, Fred
APPLICANT: Wu, Fred
APPLICANT: Pruzan, Ronald
APPLICANT: Pruzan, Ronald
APPLICANT: Weinrich, Scott L.
                                                                                                                                                                                                             US-08-770-565-5/c
; Sequence 5, Application US/08770565
; Patent No. 5846723
; GENERAL INFORMATION:
APPLICANT: Kim, Nam Woo
; APPLICANT: Wu, Fred
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 100.0
Matches 30; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 30 base pairs
                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: San Francisco
STATE: California
COUNTRY: USA
  Query Match
Best Local Similarity
Matches 30; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear MOLECULE TYPE: DNA
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US-08-770-565-8/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Andrews, William H.
APPLICANT: Andrews, William H.
APPLICANT: Feng, Junil
APPLICANT: Feng, Junil
APPLICANT: Greider, Carol
APPLICANT: Marhuenda, Maria A. B.
APPLICANT: Marhuenda, Maria A. B.
APPLICANT: Marhuenda, Waria A. B.
APPLICANT: Willeponteau, Bryant
TITLE OF INVENTION: RNA Component of Telomerase
NUMBER OF SEQUENCES: 47
CORRESPONDENCES: 47
CORRESPONDENCES ADDRESS:
ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C.
STREET: Two Militia Drive
CITY: Lexington
STRATF. Maria
                                                                                                                                                                                                                                                                                           0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: TBM PC compatible
OPERATUR SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/520,550A
FILING DATE: 29-AUG-1995
CLASSIFICATION: 435
                                                                                                                                                                                                                                     DB 1;
23;
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APPLICATION NUMBER: US 08/387,524
FILING DATE: 12-FEB-1995
FRIING DATE: 12-FEB-1995
FRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
FRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUU-1994
ATTORNEY/AGENT INFORMATION:
NAME: Granahan, Patricia
REGISTRATION NUMBER: 32,227
REBERENCE/DOCKET NUMBER: 32,227
REEFERNENCE/DOCKET NUMBER: CSHL94-05A3B
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                            77 TGCTTTTGCTCCCCGCGCGCTGTTTTCTC 106
                                                                                                                                                                                                                                     Query Match 6.7%; Score 30; DB Best Local Similarity 100.0%; Pred. No. 23; Matches 30; Conservative 0; Mismatches
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        23:
                             SEQUENCE CHARACTERISTICS:
ILENGTH: 30 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: DNA
US-08-472-802C-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: 617-00.

INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: SEQUENCE CHARACTERISTICS: SEQUENCE: 30 base pairs
        INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid
STRANDEDNESS: single
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US
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US-08-520-550A-18
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COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

APPLICATION NUMBER: US 08/72,102

FILING DATE: 07-JUL-1994

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/330,123

FILING DATE: 27-OCT-1994

ATTORNEY/AGENT INFORMATION:

NAME: SMITCH, WILLIAM M.

REGISTRATION NUMBER: 30,223

REFERENCE/DOCKET NUMBER: 15389-000820

TELECOMMONICATION INFORMATION:

TELECOMMONICATION INFORMATION:

TELECOMMONICATION INFORMATION:

TELECOMMONICATION INFORMATION:

TELEFAX: (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 23, Application US/08472802C

Patent No. 5958680
GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junii
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 44

CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CLASSIFICATION: 4.5

PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/387,524
FILING DATE: 13-FEB-1995
PRIOR APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-OCT-1994
PRIOR APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-OCT-1994
ATTORNEY, AGENT INFORMATION:
NAME: Granahan, Patricia
REGISTRATION NUMBER: SHL94-05A4
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFAX: 617-861-6240
TELEFAX: 617-861-9540
TELEFAX: 617-861-9540
INFORMATION FOR SEQ ID NO: 18:
SEQUENCE CHARACTERISTICS:
LENGTH: 30 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
STRANDEDNESS: single
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    77 TGCTTTTGCTCCCCGCGCGCTGTTTTCTC 106
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California
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 16
US-08-472-802C-23/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET: 379 Lytton Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 22, Application US/09580517
Patent No. 6320039
GENERAL INFORMATION:
         Villeponteau, Bryant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: Palo Alto
STATE: California
COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (415) 576-0300 INFORMATION FOR SEQ ID NO:
APPLICANT: Villeponteau,
APPLICANT: Feng, Junli
APPLICANT: Funk, Walter
APPLICANT: Andrews, Will
TITLE OF INVENTION: Mamm
VUMBER OF SEQUENCES: 25
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend an
                                      Feng, Junli
Funk, Walter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 30 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                           COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE:
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                                         0; Indels
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Sequence 22, Application US/0898443

Sequence 22, Application US/0898443

Sequence 22, Application US/0898443

SEMERAL INCRMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Unilia
APPLICANT: Funk, Walter
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
APPLICANT: Andrews, William H.
CITTE OF SEQUENCES: 30

CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LIP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
SCANMANDERS CONTANTON: CONTANTON CON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/998,443
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity 100.0%; Pred. No. 23; Matches 30; Conservative 0; Mismatches
                                                                                                  77 recrirrecreeceeeecererrirere 106
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Patent No. 6258535
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: DNA
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US-09-060-523-22/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY:
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Gape .; 0 FENG, Jun1i
FUNK, Walter
FUNK, Walter
TITLE OF INVENTION: HUMAN TELOMERASE
NUMBER OF SEQUENCES: 25
CORRESPONDENCES: ADDRESS:
ADDRESSEE: Townsend and Townsend Khourie and Crew 6.7%; Score 30; DB 1; Length 30; 00.0%; Pred. No. 23; Indela ADDRESSEE: Townsend and Townsend and Crew LL STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco STATE: California 77 TGCTTTTGCTCCCCGCGCGCTGTTTTCTC 106 Query Match 6.7%; Score 30; DB Best Local Similarity 100.0%; Pred. No. 23; Matches 30; Conservative 0; Mismatches 30 recririrecrecededecederarirere 1 Andrews, William H. JENTION: Mammalian Telomerase

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Best Local Similarity 100.0%; Pred. No. 23; Matches 30; Conservative 0; Mismatches
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US-09-717-828B-5/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ó
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                                                                   MEDIUM TYPE: Floppy disk
COMPUTER: IBW PC compatible
BERATURG SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
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APPLICANT: Weinrich, Scott L
APPLICANT: Atkinson III. Edward M
APPLICANT: Vasserot, Allain P
APPLICANT: Prizan, Ronald A
ITITLE OF INVENTION: A Method for Purifying Telomerase
CURRENT PRICATION NUMBER: US/09/717,828B
CURRENT FILING DATE: 1090-11-20
FRIOR APPLICATION NUMBER: 09/420,056
FRIOR APPLICATION NUMBER: 08/833,377
FRIOR APPLICATION NUMBER: 08/833,377
FRIOR FILING DATE: 1990-04-04
FRIOR FILING DATE: 1995-08-04
FRIOR FILING DATE: 1995-08-04
SRIOR APPLICATION NUMBER: 08/510,736
FRIOR APPLICATION NUMBER: 08/5
                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 08/330,123
FILING DATE: cunknown>
ATTORNEY/AGENT INFORMATION:
NAME: Smith, william M
REGISTRATION NUMBER: 30,223
REBERBNEWEY DOCKNEY: 15389-000810
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 326-2400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        77 TGCTTTTGCTCCCGCGCGCGCTGTTTTCTC 106
                                                                                                                                                                                                                       CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/580,517
FILING DATE: 25-May-2000
CLASSIFICATION: vUnknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30 TGCTTTTGCTCCCGGGGGCTGTTTTCTC 1
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SEQUENCE DESCRIPTION: SEQ ID NO: 22:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 4, Application US/09717828B Patent No. 6517834
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (415) 326-2422
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 30 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INFORMATION FOR SEQ ID NO: 22:
SEQUENCE CHARACTERISTICS:
                             COMPUTER READABLE FORM:
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: misc_feature
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US-09-717-828B-4/c
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6.7%; Score 30; DB 1; Length 30;

Query Match

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        Gaps
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; OTHER INFORMATION: Description of Artificial Sequence: Affinity Agent
US-09-717-828B-5
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        Indels
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                                                                                                                                                                                                                                                                                                                                      Sequence 5, Application US/09717828B
Fatent No. 6517834
GENERAL INFORMATION:
APPLICANT: Weinrich, Scott L
APPLICANT: Hichteeiner, Serge P
APPLICANT: Lichtseiner, Serge P
APPLICANT: Lichtseiner, Serge P
APPLICANT: Lichtseiner, Serge P
APPLICANT: Lichtseiner, Serge P
APPLICANT: PurifiedThain P
APPLICANTON NUMBER: US/09/717,828B
CURRENT FILING DATE: 1999-10-18
PRIOR FILING DATE: 1999-10-18
PRIOR FILING DATE: 1999-10-18
PRIOR PELICATION NUMBER: 08/810,736
PRIOR PELICATION NUMBER: 08/510,736
PRIOR PILING DATE: 1995-08-04
NUMBER OF SEQ ID NOS: 11
SOFTWARE: PatentIn Ver. 2.1 edited
SEQ ID NO 5
LENGTH: NAME
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GENERAL INFORMATION:
APPLICANT: Weinrich, Scott L
APPLICANT: Atkinson III, Edward M
APPLICANT: Lichtsteiner, Serge P
APPLICANT: Lichtsteiner, Serge P
APPLICANT: Lichtsteiner, Serge P
APPLICANT: Pruzan, Ronald A
TILLE REFERENCE: PurifiedTelomerase011base
CURRENT APPLICATION NUMBER: US/09/717,829A
CURRENT APPLICATION NUMBER: 09/420,056
PRIOR PELING DATE: 1999-10-18
FRIOR APPLICATION NUMBER: 08/833,377
PRIOR FILING DATE: 1999-04-04
PRIOR FILING DATE: 1999-04-04
PRIOR FILING DATE: 1995-08-04
PRIOR SPELICATION NUMBER: 08/510,736
PRIOR APPLICATION NUMBER: 08/510,736
PRIOR APPLICATION NUMBER: 08/510,736
SPRIOR APPLICATION NUMBER: 08/510,736
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                                                                      167 AAACAAAAATGTCAGCTGCTGGCCCGTTC 196
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                                                                                                                                                          30 AAACAAAAATGTCAGCTGCTGGCCCGTTC 1
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; Sequence 4, Application US/09717829A
; Patent No. 6545133
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)
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Townsend and Crew LLP

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Townsend and
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STRANDEDNESS: sing
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ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ઠે
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                   TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)
OTHER INFORMATION: Biotin S'-terminal
OTHER INFORMATION: Description of Artificial Sequence: Affinity Agent
US-09-717-829A-4
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OTHER INFORMATION: Description of Artificial Sequence: Affinity Agent
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                                                                                                                                                                                                                                 Query Match 6.7%; Score 30; DB 1; Length 30;
Best Local Similarity 100.0%; Pred. No. 23;
Matches 30; Conservative 0; Mismatches 0; Indels
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100.0%; Pred. No. 23;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Machinich, Scott L
APPLICANT: Atkinson III, Edward M
APPLICANT: Lichtereiner, Serge P
APPLICANT: Lichtereiner, Serge P
APPLICANT: Lichtereiner, Serge P
APPLICANT: Pruzan, Rohald A
TITLE OF INVENTION: A Method for Purifying Telomerase
TITLE OF INVENTION: A Method for Purifying Telomerase
CURRENT APPLICATION NUMBER: US/09/717,829A
CURRENT FILING DATE: 1000-11-20
PRIOR APPLICATION NUMBER: 08/833,377
PRIOR APPLICATION NUMBER: 08/833,377
PRIOR APPLICATION NUMBER: 08/510,736
PRIOR APPLICATION NUMBER: 08/510,736
PRIOR PILING DATE: 1995-08-04
NUMBER OF SEQ ID NOS: 11
SOFTWARE: PatentIn Ver. 2.1 edited
SEQ ID NO S
                                                                                                                                                                                                                                                                                                                                  167 AAACAAAAATGTCAGCTGCTGCCCGTTC 196
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APPLICANT: Feng, Junli
APPLICANT: Funk, Walter
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 42
                                                                                                                                                                                                                                                                                                                                                                              30 AAACAAAAATGTCAGCTGCTGGCCGTTC 1
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Patent No. 6545133
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Patent No. 6548298
GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
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Matches 30; Conservative
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US-09-057-351-22/c
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                                                                                                                                       COMPUTER: IDEN C. C. C. C. C. COMPUTER: DECADOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/057,351
FILING DATE: 08-APR.1994
CLASSIPICATION 1435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION NUMBER: US 08/472,802
FILING DATE: 07-JUN-1995
ATTORNEY/AGRY INFORMATION:
APPLICATION NUMBER: US 08/472,802
FILING DATE: 07-JUN-1995
ATTORNEY/AGRY INFORMATION:
REFERENCE/ODCKET NUMBER: 32,944
REFERENCE/ODCKET NUMBER: 32,944
REFERENCE/ODCKET NUMBER: 015389-000821US
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION SEQ ID NO: 22:
ERNGTH: 30 base pairs
WVDE: nucleic acid
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
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; OTHER INFORMATION: synthetic oligomer
US-09-903-461-2
                                                                        ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 6.73
Best Local Similarity 100.0
Matches 30; Conservative
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/note= "N = 5' biotinylated guanosine"
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23;
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ZIP: 94111-3834

ZIP: 94111-3834

COMPUTER READABLE FORM:
MEDIUW TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/833,377

FILING APPLICATION: 435
PRIOR APPLICATION NUMBER: US 08/510,736
FILING DATE: 04-APR-1997
ATPONINY/AGENT INPOMER: US 08/510,736
FILING DATE: 04-AUG-1995
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OTHER INFORMATION: /note= "oligonucleotide anti-P"
US-08-8833-377-2
                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 2, Application US/08833377
Patent No. 5965506
GENERAL INFORMATION, SCOTT L.
APPLICANT: Weinson III, Edward M.
APPLICANT: Lichtsteiner, Serge P.
APPLICANT: Vasseror, Alain P.
APPLICANT: Pruzan, Ronald A.
APPLICANT: Kealey, James T.
TITLE OF INVENTION: Purified Telomerase
NUMBER OF SEQUENCES: 15
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Crew LLP
                                                                                                                                                                                                                                   .,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: Two Embarcadero Center, Bighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                           137 CCTGCCGCCTTCCACGTTCATTCTAGAGC 166
                                                                                                                                                                                    6.7%; Score 30; DB 100.0%; Pred. No. 23; Live 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: SCORELLA, JOHN R.
REGISTRATION NUMBER: 32,944
REPRENCE/DOCKET NUMBER: 01531
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
                         SOFTWARE: PatentIn version 3.1
SEQ ID NO 5
LENGTH: 30
TYPE: DNA
CRGANIEM: HOMO Sapiens
US-10-330-872-5
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Best Local Similarity 100.(
Matches 30; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 30 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
       NUMBER OF SEQ ID NOS: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION:
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                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: Geron Corporation
APPLICANT: Weintid, Scott
APPLICANT: Weintid, Scott
APPLICANT: Weintid, Scott
APPLICANT: Lichtsteiner, Seege
APPLICANT: Lichtsteiner, Seege
APPLICANT: Lichtsteiner, Seege
APPLICANT: Using Purified Telomerase to Identify Telomerase and
TITLE OF INVENTION: Using Purified Telomerase to Identify Telomerase Activators and
TITLE OF INVENTION: Inhibitors
TITLE OF INVENTION: Inhibitors
FILE REFERENCE: 011/006C
CURRENT APPLICATION NUMBER: US/10/330,872
CURRENT FILING DATE: 1995-08-04
PRIOR FILING DATE: 1995-08-04
PRIOR FILING DATE: 1999-10-18
PRIOR FILING DATE: 1999-10-18
PRIOR FILING DATE: 2000-11-20
NUMBER OF SEQ ID NOS: 11
SSOFTWARE: PatentIn Version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Geron Corporation
APPLICANT: Geron Corporation
APPLICANT: Geron Corporation
APPLICANT: Melinrich, Scott
APPLICANT: Atkinson III, Edward
APPLICANT: Lichtsteiner, Serge
APPLICANT: Lichtsteiner, Serge
APPLICANT: Lichtsteiner, Serge
APPLICANT: Druzan, Ronald
TITLE OF INVENTION: Using Purified Telomerase to Identify Telomerase Activators and
TITLE OF INVENTION: Unibitors
FILE REFERENCE: 011/006C
CURRENT APPLICATION NUMBER: 08/510,736
FRIOR APPLICATION NUMBER: 08/510,736
FRIOR FILING DATE: 1995-08-04
FRIOR FILING DATE: 1997-04-04
FRIOR FILING DATE: 1997-04-04
FRIOR FILING DATE: 1997-04-04
FRIOR FILING DATE: 1997-04-04
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Pred. No. 23;
                            0; Indels
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100.0%; Pred. No. 2.,
                                                                   137 CCTGCCGCCTTCCACCGTTCTAGAGC 166
    Pred. No. 23;
Best Local Similarity 100.0%; Pred. No. 23; Matches 30; Conservative 0; Mismatches
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                                                                                                30 CCTGCCGCCTTCCACCGTTCATTCTAGAGC 1
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APPLICATION NUMBER: 09/717,828
FILING DATE: 2000-11-20
                                                                                                                                                                                                                            Sequence 4, Application US/10330872
Patent No. 6787133
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-10-330-872-5/c
; Sequence 5. Application US/10330872
; Patent No. 6787133
; GENERAL INFORMATION:
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Best Local Similarity 100.
Matches 30; Conservative
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ORGANISM: Homo sapiens
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US-10-330-872-4/c
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LENGTH: 30
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/note= "N = 5' biotinylated guanosine"
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MEDIUW TYPE: FI-10ppy disk
COMPUTER: 1BM PC compatible
OPERATING SYSTEM: PC-50S/MS-DOS
SOCHWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/833,377
FILING DATE: 04-APR-1997
CLASSIFICATION NUMBER: US 08/510,736
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/510,736
FILING DATE: 04-AUG-1995
ATTORNEY/AGENT INFORMATION:
NAME: Stoofella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-001110US
                                                                                                                                                                                        APPLICANT: Weinrich, Scott L.
APPLICANT: Weinrich, Scott L.
APPLICANT: Atkinson III, Edward M.
APPLICANT: Lichtsteiner, Serge P.
APPLICANT: Useserot, Alain P.
APPLICANT: Kealey, James T.
TITLE OF INVENTION: Purified Telomerase
NUMBER OF SEQUENCES: 15
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embaccadero Center, Eighth Floor
CITY: San Francisco
STATE: California
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; Sequence 3, Application US/09717828B
; Patent No. 6517834
                                                                                                            Sequence 6, Application US/08833377; Patent No. 5968506; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 6: SEQUENCE CHARACTERISTICS:
LENGTH: 30 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: modified_base LOCATION: 1
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY:
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Query Match
6.4%; Score 29; DB 1; Length 30;
Best Local Similarity 100.0%; Pred. No. 28;
Matches 29; Conservative 0; Mismatches 0; Indels
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28;
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                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Weinrich, Scott L.
APPLICANT: Weinrich, Scott L.
APPLICANT: Weinrich, Scott L.
APPLICANT: Lichteteiner, Serge P.
APPLICANT: Kaaley, James T.
TITLE OF INVENTION: Purified Telomerase
NUMBER OF SEQUENCES: 15
CORRESPONDENCE ADDRESS:
ADDRESSE: Towneend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STREET: Two Embarcadero Center, Eighth Floor
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STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STREET: Two Embarcadero Center, Eighth Floor
COMPUTER: Eldowing Application Addition Add
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                                                                                                                  43 TGTCTAACCCTAACTGAGAAGGGCGTAGG 71
                                                                                                                                                  30 TGTCTAACCCTAACTGAGAAGGGCGTAGG 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: -
LOCATION: 1..30
OTHER INFORMATION: /note= "Oligo 13"
                                                                                                                                                                                                                                                                                                              Sequence 5, Application US/08833377
Patent No. 5968506
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: modified_base
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Best Local Similarity 100.0
Matches 29; Conservative
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MOLECULE TYPE:
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NAME/KEY:
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Gaps

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Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA
Matches
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OTHER INFORMATION: Description of Artificial Sequence: Affinity Agent
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; OTHER INFORMATION: Description of Artificial Sequence: Affinity Agent
US-09-717-829A-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
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Sequence 3, Application US/09717829A

Patent No. 6545133

GENERAL INFORMATION:
APPLICANT: Atkinson III, Edward M

APPLICANT: Atkinson III, Edward M

APPLICANT: Lichteteiner, Serge P

APPLICANT: Pruzan, Romald A

TITLE OF INVENTION: A Method for Purifying Telomerase
CURRENT PRICATION NUMBER: US/09/717,829A

CURRENT APPLICATION NUMBER: 09/420,056

PRIOR PILING DATE: 1999-10-18

PRIOR FILING DATE: 1999-10-18

PRIOR PILING DATE: 1999-10-18

PRIOR PILING DATE: 1995-04-04

PRIOR PILING DATE: 1995-04-04

SRIOR PILING DATE: 1995-08-04

NUMBER OF SEO ID NOS: 11

SOFTWARE: PatentIN Ver. 2.1 edited

SEQUENCY OF TENNEY OF 
                                               APPLICANT: Meinich, Scott Land APPLICANT: Atkinson III, Edward M APPLICANT: Lichtsteiner, Serge P APPLICANT: Vesseror, Allain P APPLICANT: A Vethod for Purifying Telomerase CURRENT APPLICATION NUMBER: US/09/717,828B CURRENT PILING DATE: 2000-11-20 PRIOR PELICATION NUMBER: 08/833,377 PRIOR APPLICATION NUMBER: 08/833,377 PRIOR APPLICATION NUMBER: 08/833,377 PRIOR APPLICATION NUMBER: 08/510,736 PRIOR PILING DATE: 1995-08-04 WUMBER: OF SEQ ID NOS: 11 SOFTWARE: Patentin Ver: 2.1 edited SEQ ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1;
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Pred. No. 31;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  6.3%;
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ORGANISM: Artificial Sequence
                                   APPLICANT: Weinrich, Scott L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 30
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Best Local
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APPLICANT: Weinrich, Scott

APPLICANT: Attinson III, Edward
APPLICANT: Attinson III, Edward
APPLICANT: Lichtsteiner, Serge
APPLICANT: Lichtsteiner, Serge
APPLICANT: Lichtsteiner, Serge
APPLICANT: Pruzan, Ronald
TITLE OF INVENTION: Using Purified Telomerase to Identify Telomerase Activators and
TITLE OF INVENTION: Using Purified Telomerase to Identify Telomerase Activators and
TITLE OF INVENTION: Using Purified Telomerase to Identify Telomerase Activators and
TITLE OF INVENTION: Using Purified Telomerase to Identify Telomerase Activators and
TITLE OF INVENTION: Using Purified Date: 2002-12-24
PRIOR APPLICATION NUMBER: 08/510,736
PRIOR FILING DATE: 1997-04-04
PRIOR FILING DATE: 1997-04-04
PRIOR FILING DATE: 1999-10-18
PRIOR FILING DATE: 1999-10-18
PRIOR FILING DATE: 2000-11-20
NUMBER OF SEQ ID NOS: 11
SOFTWARRE: Patentin Version 3.1
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APPLICANT: Johns Hopkins University
APPLICANT: Johns Hopkins University
APPLICANT: Gender, Carol W.
APPLICANT: Le, Siyuan
ITILE OF INVENTION: TELOMERASE-ASSOCIATED PROTEINS
FILE REFERENCE: 07265/157001
CURRENT APPLICATION NUMBER: US/09/286,959B
CURRENT FILING DATE: 1999-04-06
PRIOR APPLICATION NUMBER: 60/080,783
PRIOR PELING DATE: 1998-04-06
NUMBER OF SEQ ID NOS: 24
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 3
LENGTH: 28
ILENGTH: 28
CORMANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 6.3%; Score 28.4; DB 1;
Best Local Similarity 96.7%; Pred. No. 31;
Matches 29; Conservative 0; Mismatches 1;
412 GAGCTGTGGGACGTGCACCCAGGACTCGGC 441
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           412 GAGCTGTGGGACGTGCACCCAGGACTCGGC 441
                                                        30 GAGCTATGGGACGTGCACCCAGGACTCGGC
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Patent No. 6300131
                                                                                                                                                                                                      Sequence 3, Application US/10330872
Patent No. 6787133
                                                                                                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: Geron Corporation
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APPLICANT: Kim, Nam Woo
APPLICANT: Kim, Nam Woo
APPLICANT: Wu, Fred
APPLICANT: Kealey, James T.
APPLICANT: Pruzan, Rother
APPLICANT: Weinrich, Scott L.
TITLE OF INVENTION: Methods for Detecting the RNA Component of
TITLE OF INVENTION: Telomerase
NUMBER OF SUMPRINCES: 26
CORRESPONDENCE: 26
CORRESPONDENCE ADDRESS:
ADDRESSEE: TOWNSEND and TOWNSEND and CREW LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-08-974-180-32/c

Sequence 32, Application US/08974180

Patent No. 6025194

Patent No. 6025194

TITLE OF INVENTION: Methods for Modulating and Identifying TITLE OF INVENTION: Callular Senescence

TITLE OF INVENTION: Callular Senescence

NUMBER OF SEQUENCES: 36

CORRESPONDENCE ADDRESS:

ADDRESSEE Geron Corporation

STREET: 230 Constitution Drive

CITY: Menlo Park

STATE: California

COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1; Length 27;
35;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUW TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/770,565
FILING DATE: 20-DEC-1996
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: TOWNSEND and TOWNSEND and CREW LLP STREET: Two Embarcadero Center, 8th Floor CITY: San Francisco STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-002300US
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-576-0200
TELEPHAX: 415-576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 6.0%; Score 27; DB Best Local Similarity 100.0%; Pred. No. 35; Matches 27; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      144 CCTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   27 CCTTCCACCGTTCATTCTAGAGCAAAC 1
                                                                                   Sequence 26, Application US/08770565; Patent No. 5846723; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO: 26: SEQUENCE CHARACTERISTICS: LENGTH: 27 base pairs TYPE: nucleic acid STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 94025
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear; MOLECULE TYPE: DNA US-08-770-565-26
                                          RESULT 37
US-08-770-565-26/c
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/note= "N = 5' biotinylated guanosine"
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                                          0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OPERATING STSTEM: PC-UDS/MD-LDG
SOFTWARE: PATCHLIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/833,377
FILING DATE: 04-APR-1997
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/510,736
FILING DATE: 04-AUG-1995
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 015389-001110US
TELEPHONE: (415) 576-0300
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 30 base pairs
TTYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                   APPLICANT: Weinrich, Scott L.
APPLICANT: Atkinson III, Edward M.
APPLICANT: Lichtsteiner, Serge P.
APPLICANT: Lichtsteiner, Serge P.
APPLICANT: Pruzan, Ronald A.
APPLICANT: Pruzan, Ronald A.
APPLICANT: Pruzan, Ronald A.
APPLICANT: Pruzan, Purified Telomerase
NUMBER OF SEQUENCES: 15
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Bighth Floor
CITY: San Francisco
STATE: California
                  Pred. No. 31;
                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            412 GAGCTGTGGGACGTGCACCCAGGACTCGG 440
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                                                                                                                    1 GCCTGGGAGGGTGGTGCCATTTTTG 28
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LOCATION: 1..30
OTHER INFORMATION: /note= "Oligo 5"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 94111.3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
100.0%; Pre
                                                                                                                                                                                                                     US-08-833-377-4/c
; Sequence 4, Application US/08833377
Patent No. 5968506
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: modified_base LOCATION: 1
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Best Local Similarity 96.6
Matches 28; Conservative
                Best Local Similarity 100.
Matches 28; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: DNA
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Gaps

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Sequence 40, Application US/08482115B
Fatent No. 5776679
GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junii
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Telomerase
TITLE OF INVENTION: Telomerase
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                       5.9%; Score 26.4; DB 1; Length 28; 96.4%; Pred. No. 41; Live 0; Mismatches 1; Indels
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STREET: Two Embacadero Center, Eighth Floor
CITY: San Francisco
STATE: California
COUNTRY: California
COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: FOODS/MS-DOS
SOFTWARE: IBM PC Compatible
OPREATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PACENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/482,115B
FILING DATE: O7-UUN-1995
CLASSIFICATION NUMBER: US 08/272,102
FILING DATE: O7-UUL-1994
PRIOR APPLICATION NUMBER: US 08/330,123
FILING DATE: Z7-OCT-1994
ATTORNEY/AGENT INFORMATION:
APPLICATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944
ATTORNEY/AGENT INFORMATION:
NAME: STOCKET NUMBER: 32,944
REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944
REFERENCE/OFCKET NUMBER: 32,944
R
                                       REFERENCE/DOCKET NUMBER: 015389-000830US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17 GCCTGGGAGGGGTGGTGGCCATTTTTG 44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 GCCTGGGAGGGTGGTGCTATTTTG 28
                                                                                            TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 31:
SEQUENCE CHARACTERISTICS:
LENGTH: 28 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                     TELECOMMUNICATION INFORMATION: TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 96.4'
Matches 27; Conservative
                                                                                                                                                                                                                                                                                                                            TOPOLOGY: linear MOLECULE TYPE: DNA
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MOLECULE TYPE:
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APPLICANT: Villeponteau, Bryant
APPLICANT: Freng, Junii
APPLICANT: Freng, Junii
APPLICANT: Freng, Junii
APPLICANT: Andrews, William H.
ITTLE OF INVENTION: Assays for the RNA Component of Human
ITTLE OF INVENTION: Telomerase
NUMBER OF SEQUENCES: 40
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                6.0%; Score 27; DB 1; Length 27; 100.0%; Pred. No. 35;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Indels
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MG-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/974,180
FILING DATE: 19-NOV-1997
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: KRATER, Kevin R.
REGISTRATION NUMBER: 32,704
REFERENCE/DOCKET NUMBER: 206
TELECOMMUNICATION INFORMATION:
TELEPRAX: (650) 473-7779
TELEPRANES: single
STRANDENESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: CALLLIANTS

ZIP: 94111-3834

ZUP: 94111-3834

COMPUTER READABLE FORM:
MEDIUW TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/482,115B

FILING DATE: 07-JUN-1995

PROME APPLICATION NUMBER: US 08/272,102

PILING DATE: 07-JUL-1994

PROM APPLICATION NUMBER: US 08/330,123

FILING DATE: 27-OCT-1994

ATTORNEY/AGENT INFORMATION:
NAME: SLOXE1la, JOHN R.
REGISTRATION NUMBER: 32,944
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; NAME/KEY: -
; LOCATION: 1..27
; CTHER INFORMATION: /note= "primer hTR445 comp"
US-08-974-180-32
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Matches 27; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: linear MOLECULE TYPE: DNA
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Sequence 23, Application US/08482115B
Sequence 23, Application US/08482115B
Sexeral INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Punk, Walter
APPLICANT: Punk, Walter
APPLICANT: Punk, Walter
APPLICANT: Punk, Walter
APPLICANT: Pank, Walter
APPLICANT: Andrews, William H.
ITILE OF INVENTION: Telomerase
NUMBER OF SEQUENCES: 40
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 5.8%; Score 26; DB 1; Length 26; Best Local Similarity 100.0%; Pred. No. 40; Matches 26; Conservative 0; Mismatches 0; Indel8
                                                                                                                                                                                                              COMPUTER READABLE FORM:

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1.25

CURRENT PPLICATION DATA:

APPLICATION NUMBER: US/08/330,123A

FILING DATE: 27-OCT-1994

CLEASSIFICATION DATA:

APPLICATION NUMBER: US 08/272,102

FILING DATE: 07-JUL-1994

ATTORNEY AGENT INFORMATION:

REFERENCE/DOCKET NUMBER: 15389-000810

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:

TELEBRONE: (415) 326-2400

TELEBRONE: (415) 326-2422

INFORMATION FOR SEQ ID NO: 23:

SEQUIENCE CHARACTERISTICS:
APPLICANT: ANDREWS, William H.
TITLE OF INVENTION: HUMAN TELOMERASE
TOWNER OF SEQUENCES: 25
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend Khourie and Crew
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              145 CTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US/08/482,115B
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ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                       STREET: 379 Lytton Avenue
CITY: Palo Alto
STATE: California
COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             26 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 26 base par
TYPE: nucleic acid
STRANDEDNESS: sing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    , MOLECULE TYPE: DNA
US-08-330-123A-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             JS-08-482-115B-23/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1; Indels
                                 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/472,802C
FILING DATE: 07-UNN-1995
CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                    APPLICANT: Villeponteau, Bryant
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junii
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION 514
PELORASIFICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M.
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 15389-000820
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 38:
ERRIGHT: 28 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Mismatches
          Pred. No. 41;
9; Mismatches
                                                                                 17 GCCTGGGAGGGGTGGTGGCCATTTTTG 44
                                                                                                             1 GCCUGGGAGGGUGGUGGCUAUUUUUG 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17 GCCTGGGAGGGGTGGTGGCCATTTTTG 44
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Sequence 23, Application US/08330123A
Sequence 23, Application US/08330123A
Pertent No. 5583016
Pertent No. 5583016
PERTENT: VILLEPROWINGU, Bryant
APPLICANT: FENG, Junii
APPLICANT: FUNK, Malter
                                                                                                                                                                                                                                                  ; Sequence 38, Application US/08472802C; Patent No. 5958680; GENERAL INFORMATION:
          64.3%;
          Best Local Similarity 64.3
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches 27; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: nucleic acid
STRANDEDNESS: sing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
                                                                                                                                                                                                      RESULT 41
US-08-472-802C-38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-08-472-802C-38
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                                                                                                                                                                                                                                                                                                0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STATE: CALITCLILIA
CCUNTRY: USA
ZIP: 9411-3834

ZIP: 9411-3834

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/660,678A

FILING DATE: 05-UDN-1996
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-CCT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 32,944

FILING DATE: 07-UUL-1994
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REFERENGE/DOCKET NUMBER: 32,944

ATTORNEY/AGENT INFORMATION:
TELEPHONE: (415) 576-0300

TELEPHONE: (415) 576-0300

TELEPHONE: (415) 576-0300

TELEPHONE: MARCTERISTICS:
LENGTH: CA base pairs

WUNDER: MUNDER: 23:
SEQUENCE CHARACTERISTICS:
LENGTH: 26 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junii
APPLICANT: Peng, Junii
APPLICANT: Punk, Walter
APPLICANT: Markews, William H.
TITLE OF INVENTION: Mammalian Telomerase
TITLE OF SECUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STREE: California
                                                                                                                                                                                                                                               Query Match 5.8%; Score 26; DB Best Local Similarity 100.0%; Pred. No. 40; Matches 26; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   145 CTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                                                                                                                 45 TCTAACCCTAACTGAGAAGGGCGTAG 70
                                                                                                                                                                                                                                                                                                                                                                                                1 rcraacccraacreagaaggggrag 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 45
US-08-660-678A-23/c
'Sequence 23, Application US/08660678A
'Patent No. 5837857
                    INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
LENGTH: 26 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: DNA
  (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 100.
Matches 26; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    linear
                                                                                                                                                                                                    US-08-482-115B-29
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  TELEFAX:
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APPLICANT: Feng, Junii
APPLICANT: Feng, Junii
APPLICANT: Funk, Walter
APPLICANT: Andrews, William H.
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Assays for the RNA Component of Human
TITLE OF INVENTION: Telomerase
NUMBER OF SEQUENCES: 40
CORRESPONDENCE ADDRESS: 40
CORRESPONDENCE ADDRESS: 40
CORRESPONDENCE ADDRESS: ADDRESSE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 26; DB 1; Length 26;
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CALP: 94111-3834
CALP: 94111-3834
CAUDUM TYPE: Floppy disk
COMPUTER: END FO compatible
COREDATIOR SYSTEM: FC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/482,115B
FILING DATE: 07-JUN-1995
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
FILING DATE: 07-JUL-1994
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-CCT-1994
ATTORNEY/AGENT INFORMATION:
NAME: SCOTELLA JOHN R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 32,944
REFERENCE/COCKET NUMBER: 015389-000830US
TELECOMMULICATION INFORMATION:
TELECOMMULICATION INFORMATION:
TELECOMMULICATION INFORMATION:
TELECOMMULICATION INFORMATION:
TELECOMMULICATION INFORMATION:
               CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UNDRER:
FILING DATE: 07-UL-1994
PRIOR APPLICATION UNDRER: US 08/272,102
PRIOR APPLICATION UNDRER: US 08/330,123
PRIOR APPLICATION UNDRER: US 08/330,123
PILING DATE: 27-OCT-1994
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION UNDRER: 32,944
REFERENCE/DOCKET WUNBER: 015389-000830US
TELEPOMMUNICATION INFORMATION:
TELEPAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 23:
SEQUENCE CHARACTERISTICS:
LENGTH: 26 base pairs
TYPE: Uncleic acid
STRANDENNES: single
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100.0%; Pred. No. *v.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 29, Application US/08482115B patent No. 5776679 GENERAL INFORMATION:
07-JUN-1995
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Best Local Similarity 100.C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: USA
ZIP: 94111-3834
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US-08-482-115B-29
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nucleic acid
EDNESS: single
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            San Francisco
                                                    California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              , MOLECULE TYPE: DNA
US-08-710-249-25
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US-09-10-249-25
US-08-10-249-25
Sequence 25, Application US/08/10249
Sequence 25, Main US/08/10249
Sequence 25, Application US/08/10249
Sequence 25
                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Wu, Fred
APPLICANT: Kealey, James T.
APPLICANT: Fealey, James T.
APPLICANT: Pruzan, Rotter
APPLICANT: Pruzan, Record L.
TITLE OF INVENTION: Methods for Detecting the RNA Component of
TITLE OF INVENTION: Telomerase
NUMBER OF SUGUENES: 26
CORRESPONDENCE ADDRESS: ADDRESSE: TOWNSEND and TOWNSEND and CREW LLP
STREET: TWO Embarcadero Center, 8th Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/770,565
FILING DATE: 20-DEC-1996
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: SLOCELIA, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/POCKET NUMBER: 015389-002300US
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-576-0200
INFORMATION FOR SEQ ID NO: 25:
SEQUENCE CHARACTERISTICS:
LENGTH: 26 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         E: Townsend and Townsend and Crew LLP
Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             60 GAAGGCCTAGGCCCCCTGCTTTTGC 85
26 CTTCCACGTTCATTCTAGAGCAAAC 1
                                                                                                                                                                                                                                    Sequence 25, Application US/08770565 Patent No. 5846723 GENERAL INFORMATION: GENERAL INFORMATION: APPLICANT: Kim, Nam Woo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.08; Pr.
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Best Local Similarity
Matches 26; Conserve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                          RESULT 46
US-08-770-565-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-08-770-565-25
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STATE: California

CONTRY: USA

CONTRES: USA

CONTRY: USA

CONTRY: USA

CONTRY: USA

CONTRY: USA

CONTRY: USA

CONTRES: USA

CONTRES: USA

CONTRY: USA

CONTRY: USA

CONTRY: USA

CONTRES: USA

CONTRES:
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Gaps
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                                                                                                                                                                                                                                            DB 1; Length 26; 40;
                                                                                                                                                                                                                                                                                         0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/472,802C
FILING DATE: 07-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junil
APPLICANT: Feng, Junil
APPLICANT: Feng, Junil
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 44
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PULLA STELLION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
FRICA PPELICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
ATTORNEY/AGENT INFORMATION:
NAME: Smith, william M.
REGISTRATION NUMBER: 30,223
REBERRENCE/DOCKET NUMBER: 15389-000820
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                            Query Match 5.8%; Score 26; DB Best Local Similarity 100.0%; Pred. No. 40; Matches 26; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                          145 CTTCCACCGTTCATTCTAGAGCAAAC 170
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; Sequence 24, Application US/08472802C
; Patent No. 5958680
; GENERAL INFORMATION:
                                   INFORMATION FOR SEQ ID NO: 19: SEQUENCE CHARACTERISTICS: LENGTH: 26 base pairs TYPE: nucleic acid STRANDEDNESS: single
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 24:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 5.8
Best Local Similarity 100.
Matches 26, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET: Two Embarcades CITY: San Francisco STATE: California COUNTRY: USA ZIP: 94111-3834 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy di
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
                                                                                                                                                                            ; TOPOLOGY: linear
US-08-485-778-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: linear MOLECULE TYPE: DNA US-08-472-802C-24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Villeponteau, Bryant
TITLE OF INVENTION: RNA COMPONENT OF TELOMERASE
NUMBER OF SEQUENCES: 45
CORRESPONDENCE ADDRESS:
ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C.
STREET: Two Militia Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: BEACHLING Misk
COMPUTER: BEACHLIN RELEASE #1.0, Version #1.30
SOCTWARE: Patentin Release #1.0, Version #1.30
SOCTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/485,778
FLING DATE: 07-15-1995
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/387,524
FLING DATE: 13-FEB-1995
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-0CT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/310,123
FILING DATE: 07-0CT-1994
ATTORNEY/AGENT INFORMATION:
                FILING DATE: 05-JAN-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 06/003,492
FILING DATE: 08-SEP-1995
ATTORNEY/AGENT INFORMATION:
NAME: SECRELIA, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-001220US
TELEPHONE: (415) 576-0200
TELEPEAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 26:
SEQUENCE CHARACTERISTICS:
LENGTH: 26 base pairs
TYPE: nucleic acid
STRANDENNES: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Greider, Carol
Marhuenda, Maria Antonia Blasco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Granahan, Patricia
REGISTRATION NUMBER: 32,227
REFERENCE/DOCKET NUMBER: CSHL94-05A4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           145 CTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CTTCCACCGTTCATTCTAGAGCAAAC 1
  APPLICATION NUMBER: US 08/583,808
FILING DATE: 05-JAN-1996
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Avilion, Ariel Athena
Feng, Junli
Funk, Walter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 19, Application US/08485778
Patent No. 5876979
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 5.8
Best Local Similarity 100.
Matches 26; Conservative
                                                                                                                                                                                                                                                                                                                                                                                            linear
: DNA
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APPLICANT: Andrews
APPLICANT: Avilion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Lexington
                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: 1i
MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SS
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US-08-485-778-19/c
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APPLICANT:
APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET:
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US-08-974-180-33
; Sequence 33, Application US/08974180
; Patent No. 6025194
; Patent No. 6025194
; GENERAL INFORMATION:
; TITLE OF INVENTION: Methods for Modulating and Identifying
; TITLE OF INVENTION: Callular Senescence
; TITLE OF INVENTION: Callular Senescence
; MUMBER OF SEQUENCES: 36
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Geron Corporation
; STREET: 230 Constitution Drive
; CITY: Menlo Park
; STATE: California
; COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5.8%; Score 26; DB 1; Length 26; 100.0%; Pred. No. 40;
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TITLE OF INVENTION: RNA Component of Telomerage NUMBER OF SEQUENCES: 47
CORRESPONDENCE ADDRESS:
ADDRESSEB: Hamilton, Brook, Smith & Reynolds, P.C. STREET: Two Militia Drive
CITY: Lexington
STATE: US
STATE: US
                                                                                                                                                                                                                                                                                        CAMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: EN PC Compatible
COMPUTER: EN PC Compatible
COMPUTER: EN PC Compatible
COMPUTER: EN PC COMPOSIME
COMPUTER: EN PC COMPOSIME
COPERATION SYSTEM: PC-DCS/MS-DOS
SOFTWARE: PATENTION DATA:
APPLICATION NUMBER: US/08/520,550A
FILING DATE: 29-ANG-1995
RION APPLICATION DATA:
APPLICATION NUMBER: US 08/387,524
FILING DATE: 13-FEB-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION DATA:
APPLICATION DAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 94025
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Granahan, Parricia
REGISTRATION NUMBER: 32,227
REGISTRATION NUMBER: 32,227
RELECOMMUNICATION INDORMATION:
TELEPHONE: 617-861-6240
INFORMATION FOR SEQ ID NO: 19:
SEQUENCE CHARACTERISTICS:
LENGTH: 26 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ilarity 100.0%; Pred. No. 40;
Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           145 CTTCCACCGTTCATTCTAGAGCAAAC 170
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
Matches 26; Conserv
                                                                                                                                                                                                                                                                       02173
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US-08-520-550A-19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IMP FO Compatible

SOFTWARE: Patentin Release #1.0, Version #1.30

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/472,802C

FILING DATE: 07-JUL-1994

PRIOR APPLICATION NUMBER: US 08/330,123

FILING DATE: 27-OCT-1994

ATTORNEY/AGENT INFORMATION:

NAME: SMAIT, William M.

REGISTRATION NUMBER: 15389-000820

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                     Sequence 30. Application US/08472802C
| Sequence 30. Application US/08472802C
| GENERAL INFORMATION:
| APPLICANT: Villeponteau, Bryant
| APPLICANT: Feng, Junia H. TITLE OF INVENTION: Mammalian Telomerase
| NUMBER OF SEQUENCES: 44
| CORRESPONDENCE ADDRESS:
| ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  45 TCTAACCCTAACTGAGAGGGCGTAG 70
                                         26 criccaccericaricrasascaaec 1
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APPLICANT: Andrews, William H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Greider, Carol
Marhuenda, Maria A. B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Andrews, William H.
Avilion, Ariel A.
Feng, Junli
Funk, Walter
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE: (415) 576-0200
TELEPAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 30: SEQUENCE CHARACTERISTICS:
LENGTH: 26 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 100.0
werches 26; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 52
US-08-520-550A-19/c
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US-08-472-802C-30
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APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
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Gaps

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                                                                                                                                                                                                                                                                                      DB 1; Length 26; 40;
                                                                                                                                                                                                                                                                                                                                             0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STATE: CALILORINA
COUNTER: CALILORINA
ZIP: 9411-3834
COMPUTER READABLE FORM:
MEDIUW TYPE: F1-Doppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
GURENTY APPLICATION DATA:
APPLICATION NUMBER: US/08/974,549A
FILING DATE: 19-NOV-1997
CLASSIFICATION DATA:
APPLICATION NUMBER: US/08/974,643
FILING DATE: 19-NOV-1997
PILING DATE: 18-APR-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/844,419
FILING DATE: 18-APR-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/846,017
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/851,843
FILING DATE: 06-MA'-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/851,843
FILING DATE: 06-MA'-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/851,843
FILING DATE: 06-MA'-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/854,050
PRIOR APPLICATION NUMBER: US/08/854,050
                                                                                                                                                                                                                                                                             Query Match
5.8%; Score 26; DB
Best Local Similarity 100.0%; Pred. No. 40;
Matches 26; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                      145 CTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                                                                                                                                                                                             26 CTTCCACCGTTCATTCTAGAGCAAAC 1
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/912,951
FILING DATE: 14-AUG-1997
PRIOR APPLICATION DATE:
APPLICATION NUMBER: US 08/915,503
FILING DATE: 14-AUG-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 597, Application US/08974549A Patent No. 6166178 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 09-MAY-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/911,312
        TELEFAX: (415) 576-0300

INFORMATION FOR SEQ ID NO: 23:
SEQUENCE CHARACTERISTICS:
LENGTH: 26 base pairs
TYPE: mucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
NOLECULE TYPE: DNA
US-08-998-443-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 55
US-08-974-549A-597
                                                                                                                                                                                                                                                                                Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 23, Application US/08998443
; Sequence 23, Application US/08998443
; Patent No. 6054575
; GENERAL INFORMATION:
; APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junil
; APPLICANT: Funk, Walter
; APPLICANT: Andrews, William H.
: TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/08/998,443
FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY: -
LOCATION: 1..26

COTHER INFORMATION: /note= "primer hTR S328"
US-08-974-180-33
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APPLICATION NUMBER: US/08/660,678
FILING DATE: 05-JUN-1996
APPLICATION NUMBER: US/08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944
REFECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                306 TIGGGCTCTGTCAGCCGCGGGTCTCT 331
APPLICATION NUMBER: US/08/974,180
                                                                                          NAME: Katter, Kevin R.
REGISTRATION NUMBER: 32,704
REFRENCE/DOCKET NUMBER: 206
TELECOMMUNICATION INPORMATION:
TELEFROME (650) 473-8654
INPORMATION POR SEQ ID NO: 33:
SEQUENCE CHARACTERISTICS:
LENGTH: 26 base pairs
TYRE: nucleic acid
STRANDENESS: single
                           FILING DATE: 19-NOV-1997
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-08-998-443-23/c
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Query Match 5.8%; Score 26; DB 1; Length 26; Best Local Similarity 100.0%; Pred. No. 40; Matches 26; Conservative 0; Mismatches 0; Indels
PRIOR APPLICATION DATA:

APPLICATION DATA:
APPLICATION NUMBER: US 08/851,843
FILING DATE: 06-MAY-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/854,050
FILING DATE: 09-MAY-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/911,312
FILING DATE: 14-AUG-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/912,951
FILING DATE: 14-AUG-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/915,503
FILING DATE: 14-AUG-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/US97/17618
FILING DATE: 01-0CT-1997
PRIOR APPLICATION NUMBER: WO PCT/US97/17885
FILING DATE: 01-0CT-1997
ATTORNEY/AGENT INFORMATION:
NAME: Apple, Randcalph Ted
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 36,429
RELEPAX: (415) 576-0300
SEQUENCE CHARACTERICS:
FUNCTURED DATE: 0200
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 23, Application US/09060523
Fatent No. 6258535
GENERAL INFORMATION:
APPLICANT: Feng, Junia
APPLICANT: Feng, Junia
APPLICANT: Feng, Junia
APPLICANT: For Malter
APPLICANT: Andrews, William H.
TILLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 25
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; LOCATION: 1..26
; OTHER INFORMATION: /note= "R3c primer"
US-08-974-549A-598
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    145 CTTCCACCGTTCATTCTAGAGCAAAC 170
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COUNTRY: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                JS-09-060-523-23/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: Cech, Thomas R.
APPLICANT: Linguer, Joachim
APPLICANT: Nakamura, Toru
APPLICANT: Nakamura, Toru
APPLICANT: Morin, Gregg B.
APPLICANT: Harley, Calvin B.
APPLICANT: Andrews, William H.
TITLE OF INFORTION: Human Telomerase Catalytic Subunit
NUMBER OF SEQUENCES: 727
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: San Prancisco
CITY: California
CONTRY: USA
ZIP: 94111-3834
ZIP: 94111-3834
ZOMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
APPLICATION NUMBER: US/08/974,549A
FILING DATE: 01-0CT-1996
PRIOR APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
PRIOR APPLICATION NUMBER: US 08/844,419
FILING DATE: US 08/844,419
FILING DATE: US 08/846,017
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Two Embarcadero Center, Eighth Floor
PRIOR APPLICATION DATA:

APPLICATION NUMBER: WO PCT/US97/17618
FILING DATE: 01-0CT-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/US97/17885
FILING DATE: 01-0CT-1997
ATTORNEY/AGENT INFORMATION:
NAME: Apple, Randolph Ted
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 36,429
RECECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 597:
SEQUENCE CHARACTERISTICS:
LENGTH 25 DASE PAIRS
LENGTH 26 DASE PAIRS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: -
LOCATION: 1..26
OTHER INFORMATION: /note= "F3b primer"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  45 TCTAACCCTAACTGAGAGGGCGTAG 70
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2008-974-549A-59B/c
; Sequence 598, Application US/08974549A
; Patent No. 6166178
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend a
STREET: Two Embarcade
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-08-974-549A-597
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US-09-220-157A-26/C

Sequence 26, Application US/09220157A

Sequence 26, Application US/0922015A

Sequence 26, Application U
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APPLICATION NUMBER: US/09/220,157A
                                           FILING DATE: 08-SEP-1995
ATORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-001220US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELERAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 25:
SEQUENCE CHARACTERISTICS:
LENGTH: 26 base pairs
TYPE: nucleic acid
STRANDEDNES: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         015389-001220US
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CLASSIFICATION:
PRIOR APPLICATION:
APPLICATION NUMBER: US/08/710,249
FILING DATE: 13-SEP-1996
APPLICATION NUMBER: US 08/583,808
FILING DATE: 05-JAN-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/003,492
FILING DATE: 06-SEP-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               45 TCTAACCCTAACTGAGAAGGGCGTAG 70
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               US 60/003,492
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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REGISTRATION NUMBER: 32,949
REFERENCE/DOCKET NUMBER: 01531
TELECOMMUNICATION INFORMATION:
TELEPAX: (415) 576-0200
INFORMATION FOR SEQ ID NO: 26:
SEQUENCE CHARACTERISTICS:
               APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; MOLECULE TYPE: DNA
US-09-220-157A-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       linear
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APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junil
APPLICANT: Andrews, William H.
APPLICANT: Andrews, William H.
APPLICANT: Andrews, William H.
APPLICANT: Adams, Robert R.
TITLE OF INVENTION: Methods and Reagents for Regulating
TITLE OF INVENTION: Telomere Length and Telomerase Activity
NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 5.8%; Score 26; DB 1; Length 26; Best Local Similarity 100.0%; Pred. No. 40; Matches 26; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATIOR SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/220,157A
FILING DATE:
CLASSIFICATION NUMBER: US/08/710,249
FILING DATE: 13-52P-1996
FILING DATE: 10-52P-1996
FILING DATE: 05-JAN-1996
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/060,523
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STREET: Two Embarcadero Center, Eighth Floor
SITY: San Francisco
STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   015389-000813US
                                                                                                               CLASSIETCATION. 536
CLASSIETCATION. 536
PRIOR APPLICATION NUMBER: US 08/660,678
FILING DATE: 05-UNU-1996
PRIOR APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT 1994
PRIOR APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-UL-1994
PRIOR APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-UL-1994
ATTORNEY APPLICATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-0006
TELECOMMUNICATION INFORMATION:
TYPE: INDOLESIC ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          145 CTTCCACCGTTCATTCTAGAGCAAAC 170
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                                                               UMBER: US/09/060,523
14-APR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 25, Application US/09220157A Patent No. 6300110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
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US-09-060-523-23
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US-09-220-157A-25
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DB 1;
40;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear
MOLECULE TYPE: DNA
SEQUENCE DESCRIPTION: SEQ ID NO: 23:
US-09-580-517-23
                               CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/330,123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         145 CTTCCACCGTTCATTCTAGAGCAAAC 170
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FILING DATE: 25-May-2000
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 26 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cech, Thomas R.
Lingner, Joachim
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Patent No. 6120039
GENERAL INFORMATION:
APPLICANT: VILLEPONTEAU, Bryant
FENG, Junli
FUNK, Walter
ADDRESSEE: SQUENCES: 25
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend Khourie and Crew
STREET: 379 Lytton Avenue
                                                                                                                                                                                                     5.8%; Score 26; DB 1; Length 26;
100.0%; Pred. No. 40;
tive 0; Mismatches 0; Indels
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFFWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 60
US-09-286-9598-4/C
Sequence 4, Application US/09286959B
Patent No. 6300131
GENERAL INFORMATION:
APPLICANT: Johns Hopkins University
APPLICANT: Johns Hopkins University
APPLICANT: Le, Siyuan
TILE OF INVENTION: TELOMERASE-ASSOCIATED PROTEINS
FILE REFERENCE: 07265/157001
CURRENT APPLICATION NUMBER: US/09/286,959B
CURRENT FILING DATE: 1999-04-06
PRIOR APPLICATION NUMBER: 60/080,783
PRIOR FILING DATE: 1998-04-06
NUMBER OF SEQ ID NOS: 24
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 4
LENGTH: 26
TURNET TO THE TO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 5.8%; Score 26; DB Best Local Similarity 100.0%; Pred. No. 40, Matches 26; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                           145 CTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 145 CTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                26 criccaccerrcaricrasascaasca
                                                                                                                                                                                                                                                                                                                                                                                                CTTCCACCGTTCATTCTAGAGCAAAC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 94301
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: Palo Alto
STATE: California
COUNTRY: US
                                                                                                                                                                                                                                        Best Local Similarity 100.
Matches 26; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: Primer US-09-286-959B-4
                                          single
TYPE: nucleic acid
STRANDEDNESS: singl
TOPOLOGY: linear
MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-09-580-517-23/c
                                                                                                             ; MOLECULE TYPE
US-09-220-157A-26
                                                                                                                                                                                                            Query Match
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DIAGNOSTIC AND Gaps ö APPLICANT: Linguer, Joachim
APPLICANT: Nakamura, Toru
APPLICANT: Nakamura, Toru
APPLICANT: Nakamura, Toru
APPLICANT: Morin, Greeg B.
APPLICANT: Harley, Calvin
TITLE OF INVENTION: THERAPEUTIC METHODS
NUMBER OF SEQUENCES: 335
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, 8th Floor
CITY: San Francisco
STATE: California
CCUNTRY: United States of America Length 26; Indels COMPUTER READABLE FORM:
MEDIUM TYPE: Floppadible
COMPUTER: IBM PC compatible
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/912,951
FLING DATE: 14-AUG-1997
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/854,050
FLING APPLICATION 1435
PRIOR APPLICATION NUMBER: US 08/851,843
FLING DATE: 25-APR-1997
FILING DATE: 25-APR-1997 ATTORNEY/AGENT INFORMATION:

ATTORNEY/AGENT INFORMATION:

NAME: SAILT, William M
REGISTRATION NUMBER: 30,223

REFERENCE/DOCKET NUMBER: 15389-000810

TELEPHONE: (415) 326-2420

INFORMATION FOR SEQ ID NO: 23:

SEQUENCE CHARACTERISTICS:

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5.8%; Score 26; DB 1; Length 26; 100.0%; Pred. No. 40; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         result 64
US-057-351-23/C

18-09-057-351-23/C

18-09-1057-351

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APPLICATION NUMBER: US/09/057,351
FILING DATE: 08-APR-1994
CLASSIFICATION NUMBER: US/09/057,351
FILING DATE: 08-APR-1994
PRIOR APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
PRIOR APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION DATE: US/08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION NUMBER: US 08/472,802
FILING DATE: 07-JUN-1995
                                     145 CTTCCACCGTTCATTCTAGAGCAAAC 170
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity 100.
Matches 26; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE: DNA
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Patent No. 6475789

GENERAL INFORMATION:
APPLICANT: Cingner, Joachim
APPLICANT: Chapman, Karen B.
APPLICANT: Chapman, Karen B.
APPLICANT: Andrews, William H.
APPLICANT: Andrews, William H.
APPLICANT: Harley, Calvin
APPLICANT: TITLE OF INVENTION: THERAPEUTIC METHODS
TITLE OF INVENTION: THERAPEUTIC METHODS
NUMBER OF SEQUENCE: 335
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
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100.0%; Pred. No. 40;
Live 0; Mismatches 0; Indels
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CITY: San Francisco
STATE: California
COUTRY: United States of America
ZIP: 94111
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/912,951
FILING DATE: 14-AUG-1997
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/854,050
FILING DATE: 09-MAX-1997
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/851,843
FILING DATE: 06-MAY-1997
CLASSIFICATION: 435
FILING DATE: US 08/851,843
FILING DATE: US 08/851,843
FILING DATE: US 08/MSY-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Crew LLP
STREET: Two Embarcadero Center, 8th Floor
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION DATA:
FILING DATE: US 08/844,419
FILING DATE: US 08/844,419
CLASSIFICATION: 435
PRIOR APPLICATION 1435
PRIOR APPLICATION NUMBER: US 08/724,643
FILING DATE: 01-0CT-1996
CLASSIFICATION: 435
ATTORNEY, AGENT INFORMATION:
NAME: Apple, Randolph T.
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002600US
TELEFARATION: (415) 576-0200
TELEFARA: (415) 576-0300
TELEFARA: (415) 576-0300
TELEFARA: (415) 576-0300
TELEFAR: (415) 576-0300
TELERATICON FOR SEQ ID NO: 311:
SEQUENCE CHARACTERISTICS:
LENGTH: 26 base pairs
TYPE: UNCleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        45 TCTAACCCTAACTGAGAAGGGCGTAG 70
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Best Local Similarity 100.0
Matches 26, Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: DNA US-08-912-951-311
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 63
US-08-912-951-312/c
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                                                                                                                                                                                                                                                                                    Sequence 4, Application US/09633573

Sequence 4, Application US/09633573

Patent No. 6607898

GENERAL INFORMATION:
APPLICANT: Kopreski, Michael S.
APPLICANT: Gocke, Christopher D.
TITLE OF INVENTION: Telomerase-associated RNA in Plasma or Serum
FILE REFERENCE: 00-1328

CURRENT APPLICATION NUMBER: US/09/653,573

CURRENT FILING DATE: 2000-08-31

PRIOR FILING DATE: 2000-08-31

PRIOR FILING DATE: 2000-08-31

NUMBER OF SEQ ID NOS: 7

SOFTWARE: Patentin Ver. 2.1

SEQ ID NO 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Kopreski, Michael S.
APPLICANT: Kopreski, Michael S.
APPLICANT: Gocke, Christopher D.
TITLE OF INVENTION: Method for Detection of hTR and hTERT
TITLE OF INVENTION: Telomerase-associated RNA in Plasma or Serum
FILE REFERENCE: 00-1328
CURRENT APPLICATION NUMBER: US/09/653,573
CURRENT APPLICATION NUMBER: 09/653,573
PRIOR FILING DATE: 2000-08-31
NUMBER OF SEQ ID NOS: 7
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 5
LENGTH: 26
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                                                                                                                  0; Indele
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                                                                         Query Match

5.8%; Score 26; DB

Best Local Similarity 100.0%; Pred. No. 40;

Matches 26; Conservative 0; Mismatches
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ilarity 100.0%; Pred. No.
Conservative 0; Mismatci
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                                                                                                                                                         45 TCTAACCCTAACTGAGAAGGGCGTAG 70
                                                                                                                                                                              1 TCTAACCCTAACTGAGAAGGGCGTAG 26
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; Sequence 5. Application US/09653573
; Patent No. 6607998
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 100.0
Matches 26; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; ORGANISM: Homo sapiens US-09-653-573-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-653-573-5
linear
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Best Local Similarity
Matches 26; Conserv
TOPOLOGY: 11
MOLECULE TYPE:
                                      US-09-057-351-30
                                                                                                                                                                                                                                                     RESULT 66
US-09-653-573-4
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100.0%; Pred. No. 40;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPIETS: USA

ZIP: 94111-3834

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
COMPOTER: IBM PC COMPATIBLE
COMPOTER: BEATHLINE STEEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/057,351
FILING DATE: 08-APR-1994
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/22,102
FILING DATE: 07-JUL-1994
PRIOR APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION NUMBER: US 08/330,123
FILING DATE: 07-JUN-1995
APPLICATION NUMBER: US 08/472,802
FILING DATE: 07-JUN-1995
ATTORNEY AGENT INFORMATION:
NAME: SCOTELLA, JOHN R.
REFERENCE/DOCKET NUMBER: 015389-000821US
TELECOMMUNICATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-000821US
        ATTOWNEY AGENT INFORMATION:
NAME: SCORELLA, John R.
REGISTRATION NUMBER: 32,944
REPERENCE/DOCKET NUMBER: 015389-000821US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0300
INFORMATION POR SEQ ID NO: 23:
SEQUENCE CHARACTERISTICS:
LENGTH: 26 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                            145 CTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                                                                                                                                                                                       26 CTTCCACCGTTCATTCTAGAGCAAAC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 30:
SEQUENCE CHARACTERISTICS:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 100.
Matches 26; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 26 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                               TOPOLOGY: linear
MOLECULE TYPE: DNA
US-09-057-351-23
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CUGURIER READABLE FORM:

MEDLUM TYPE: R10PQY disk

COMPUTER: F10PQY disk

MEDLUM TYPE: F10PQY disk

COMPUTER: 19 PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFFWARR: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/402,181B

FILING DATE: 2-Sep-1997

APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 08/844,419

FILING DATE: 10-OCT-1996
APPLICATION NUMBER: US 08/846,017

FILING DATE: 25-APR-1997
APPLICATION NUMBER: US 08/851,843

FILING DATE: 09-MAY-1997
APPLICATION NUMBER: US 08/911,312

FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/912,951

FILING DATE: 14-AUG-1997
APPLICATION NUMBER: WO PCT/US97/17885
APPLICATION NUMBER: WO PCT/US97/17885
                                                                                                                                                                                                                Morin, Calvin B.
Harley, Calvin B.
Andrews, William H.
TITLE OF INVENTION: Human Telomerase Catalytic Subunit
NUMBER OF SEQUENCES: 633
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 015389-002620US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LOCATION: 1..26

SEQUENCE DESCRIPTION: SEQ ID NO: 597:
US-09-402-181B-597
                                                    Sequence 597, Application US/09402181B
Patent No. 6610839
GENERAL INFORMATION:
APPLICANT: Cech, Thomas R.
Lingner, Joachim
Nakamura, Toru
Chapman, Raren B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Ausenhus, Scott L.
REGISTRATION NUMBER: 42,271
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 597:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION TELEPHONE: (415) 576-020
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 26 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY:
RESULT 68
US-09-402-181B-597
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5.8%; Score 26; DB 1; Length 26;

Query Match

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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  Andrews, William H.
TITLE OF INVENTION: Human Telomerase Catalytic Subunit
NUMBER OF SEQUENCES: 633
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATE:
PILING DATE: 29-Sep-1997
CLASSIFICATION: «Unknown»
                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION: cub.1997
CLASSIFICATION: cub.1997
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 08/724,643
FILING DATE: 01-0CT-1996
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/81,843
FILING DATE: 25-APR-1997
APPLICATION NUMBER: US 08/81,843
FILING DATE: 06-MAY-1997
APPLICATION NUMBER: US 08/911,312
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/911,312
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/912,951
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/912,951
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/915,503
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/915,503
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/915,503
FILING DATE: 01-0CT-1997
TUORNEY/AGENT INFORMATION.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Ausenhus, Scott L.
REGISTRATION NUMBER: 42,271
REFERENCE POCKET NUMBER: 015389-002620US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
                          0;
  Pred. No. 40;
; Mismatches
                                                                    70
                                                                                                                   1 rcraacccraacreagaagggggrag 26
                                                                    45 TCTAACCCTAACTGAGAAGGGCGTAG
                                                                                                                                                                                                             US-09-402-181B-598/c; Sequence 598, Application US/09402181B
Patent No. 6610839; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                 APPLICANT: Cech, Thomas R.
Lingner, Joachim
Nakamura, Toru
Chapman, Karen B.
Morin, Gregg B.
Harley, Calvin B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             576-0300
Best Local Similarity 100.0%; P
Matches 26; Conservative 0;
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TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO: 598:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (415)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY:
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Gaps
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Harley, Calvin B.
Andrews, William H.
TITLE OF INVENTION: Human Telomerase Catalytic Subunit
NUMBER OF SEQUENCES: 727
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                      Length 26;
                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                  NAME/KEY: -
LOCATION: 1..26
OTHER INFORMATION: /note= "F3b primer"
SEQUENCE DESCRIPTION: SEQ ID NO: 597:
                                                                                                                                                                                                                                                                                                                                                                                      DB 1;
40;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READBLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Vers
SOFTWARE: PATENTIN ROLAR:
APPLICATION NUMBER: US/09/721,456
FILING DATE: 22-No. 6617110-2000
CLASSIFICATION NUMBER: US/09/721,456
APPLICATION NUMBER: US/08/974,549A
FILING DATE: 19-NOV-1997
APPLICATION NUMBER: US 08/724,643
FILING DATE: 19-NOV-1997
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/846,017
FILING DATE: 25-APR-1997
APPLICATION NUMBER: US 08/846,017
FILING DATE: 25-APR-1997
APPLICATION NUMBER: US 08/846,017
FILING DATE: 25-APR-1997
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V 100.0%; Pred. No. ...
O; Mismatches
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APPLICATION NUMBER: US 08/911,312
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/915,503
FILING DATE: 14-AUG-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US 08/854,050
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TCTAACCCTAACTGAGAAGGGCGTAG 70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 rcraacccraacrgagaagggggrag 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-721-456-598/c
; Sequence 598, Application US/09721456
; Patent No. 6617110
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 06-MAY-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 14-AUG-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Cech, Thomas R.
Lingner, Joachim
Nakamura, Toru
Chapman, Karen B.
                                                                                              LENGTH: 26 base pairs
                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: single
                                            INFORMATION FOR SEQ ID NO: 59'
SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STATE: California COUNTRY: USA
                                                                                                                                                                    TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 26; Conserv
                                                                                                                                                                                                                                          NAME/KEY:
                        TELEFAX:
                                                                                                                                                                                                                                                                                                                                         US-09-721-456-597
                                                                                                                                                                                                                      FEATURE
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                                                                                                                                                                                                Gaps
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ZIP: 9411-3834

COMPUTER READABLE FORM:

COMPUTER: IBM PC compatible

COMPUTER: IBM PC COMPATIBLE

COMPUTER: DAGNON NOW.

CURRENT APPLICATION DATA:

APPLICATION DATA:

RILING DATE: 22-No. 6617110-2000

CLASSIFICATION: CUNKNOWN>

PRIOR APPLICATION NUMBER: US/08/974,549A

FILING DATE: 19-NOV-1997

APPLICATION NUMBER: US 08/744,643

FILING DATE: 10-OCT-1996

APPLICATION NUMBER: US 08/844,419

FILING DATE: 10-OCT-1996

APPLICATION NUMBER: US 08/846,017

FILING DATE: 18-APR-1997

APPLICATION NUMBER: US 08/846,017

FILING DATE: 21-APR-1997

APPLICATION NUMBER: US 08/846,017

FILING DATE: 21-APR-1997

APPLICATION NUMBER: US 08/846,017

FILING DATE: 21-APR-1997

APPLICATION NUMBER: US 08/851,843
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Chapman, faren B.
Morin, Gregg B.
Harley, Calvin B.
Andrews, William H.
TITLE OF INVENTION: Human Telomerase Catalytic Subunit NUMBER OF SEQUENCES: 727
CORRESPONDENCE ADDRESS:
                                                                                                                                              Length 26;
                                                                                                                                                                                           0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Townsend and Townsend and Crew LL STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Apple, Randolph Ted
REGISTRATION NUMBER: 36,429
REGISTRATION INVEST: 015389-002610US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 06-MAY-1997
APPLICATION NUMBER: US 08/854,050
FILING DATE: 09-MAY-1997
APPLICATION NUMBER: US 08/911,312
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/912,951
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/915,503
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: WO PCT/US97/17618
FILING DATE: 01-OCT-1997
APPLICATION NUMBER: WO PCT/US97/17885
FILING DATE: 01-OCT-1997
APPLICATION NUMBER: WO PCT/US97/17885
FILING DATE: 01-OCT-1997
LOCATION: 1..26

COTHER INFORMATION: /note= "R3c primer"

SEQUENCE DESCRIPTION: SEQ ID NO: 598:
US-09-402-181B-598
                                                                                                                                           Query Match 5.8%; Score 26; DB 1; Best Local Similarity 100.0%; Pred. No. 40; Matches 26; Conservative 0; Mismatches
                                                                                                                                                                                                                                          145 CTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                                        CTTCCACCGTTCATTCTAGAGCAAAC 1
                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 597, Application US/09721456
Patent No. 6617110
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cech, Thomas R.
Lingner, Joachim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nakamura, Toru
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Cech,
                                                                                                                                                                                                                                                                                                                                                                                    US-09-721-456-597
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MOLECULE TYPE: RNA (genomic)
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Best Local Similarity 72.03
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid
STRANDEDNESS: single
nucleic acid
                                         linear
                                                                               DESCRIPTION:
DESCRIPTION:
DESCRIPTION:
DESCRIPTION:
DESCRIPTION:
US-08-630-019A-26
TYPE: nucleic
STRANDEDNESS:
TOPOLOGY: lin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-08-630-019A-36
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INPORMATION:
APPLICANT: Shay, Jerry W.
APPLICANT: Wright, Woodring E.
APPLICANT: Piatyszek, Mieczyslaw A.
APPLICANT: Piatyszek, Mieczyslaw A.
APPLICANT: Ocrey, David
APPLICANT: Ocrey, David
TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 46
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  5.8%; Score 26; DB 1; Length 26;
100.0%; Pred. No. 40;
cive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/630,019A
FILING DATE: 09-UUN-1996
CLASSIFICATION: 536
                                                         ATTORANGE AND COLT-1997
ATTORNEY/AGENT INFORMATION:
NAME: APPID - Randolph Ted
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002610US
TELEPHONE: (415) 576-0200
TELEPHONE: (415) 576-0300
INFORMATION FOR ENG ID NO: 598:
SEQUENCE CHARACTERISTICS:
LENGTH: 26 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note= "R3c primer"
                      FILING DATE: 01-OCT-1997
APPLICATION NUMBER: WO PCT/US97/17885
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 015389-001600US
TELEPHONE: (415) 576-0200
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 26: SEQUENCE CHARACTERISTICS:
LENGTH: 25 base pairs
    APPLICATION NUMBER: WO PCT/US97/17618
                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: - LOCATION: 1..26
COTHER INFORMATION: /note= "R3C | SEQUENCE DESCRIPTION: SEQ ID NO: 598: US-09-721-456-598
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      145 CTTCCACGGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   26 CTTCCACCGTTCATTCTAGAGCAAAC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 26, Application US/08630019A Patent No. 6015710
                                                                                                                                                                                                                                                                                                   TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear
MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 100.
Matches 26; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: Two Embarcad
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    USA
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US-08-630-019A-26
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/desc = "peptide nucleic acid (PNA), where (deoxy/ribose-phosphate linkages are replaced by N+ (2-aminoethyl)glycine units linked to nucleotide bases via glycine amino nitrogen through a methylenecarbonyl linker"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 36, Application US/08630019A
Fatent No. 6015710
GENERAL INFORMATION:
APPLICANT: Shay, Jerry W.
APPLICANT: Wright, Woodring E.
APPLICANT: Piatyszek, Mieczyslaw A.
APPLICANT: Ocrey, David
APPLICANT: No. 6015710ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by
TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEGUENCES:
ADDRESSEE: Townsend and Crew LLP
ADDRESSEE: Townsend and Crew LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 25; DB 1; Length 25; Pred. No. 45;
                                                                                                                                                                                    Query Match 5.5%; Score 25; DB 1; Length 25; Best Local Similarity 72.0%; Pred. No. 45; Matches 18; Conservative 7; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC COMPATIBLE
COMPUTER: PEN PC COMPATIBLE
COMPUTER: DEAD FOR THE PC-DOS/MS-DOS
SOFTWARE: PATENTIN RELEASE #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/630,019A
FILING DATE: 09-UNN-1996
CLASSIFICATION: S36
ATTORNEY/AGENT INFORMATION:
NAME: SCOTELLA, JOHN R.
REFERENCE/POCKET NUMBER: 015389-001600US
TELLEPHONE: (415) 576-0300
TELLEPHONE: CHARACTERISTICS:
LENGTH: 25 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                7; Mismatches
                                                                                                                                                                                                                                                                                                 41 TTTGTCTAACCCTAACTGAGAAGGG 65
                                                                                                                                                                                                                                                                                                                                   1 UUUGUCUAACCCUAACUGAGAAGGG 25
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MOLECULE TYPE: other nucleic acid
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Gaps
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Sequence 28, Unit, Walter
APPLICANT: Villeport Walter
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Assays for the RNA Component of Human
TITLE OF INVENTION: Assays for the RNA Component of Human
TITLE OF INVENTION: Telomerase
NUMBER OF SEQUENCES: 40
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 25;
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Compatible
COMPUTER: Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/482,115B
FILING DATE: 07-JUN-1995
CLASSIFICATION: 435
                                                                        ZONE SALE SALE SORM:

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/349,532
                                                                                                                                                                                                                                                                                                                APPLICATION NOMBER: US/09/349,532
CLASSIFICATION DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/638,545
FILING DATE: 09-APR-1997
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-001610US
TELEFERNOR: (415) 576-0200
TELEFERNOR: (415) 576-0300
TELEFERX: (415) 576-0300
TELEFEX: (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 5.5%; Score 25; DB 1; Best Local Similarity 72.0%; Pred. No. 45; Matches 18; Conservative 7; Mismatches (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                41 TTTGTCTAACCCTAACTGAGAAGGG 65
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APPLICATION NUMBER: US 08/272,102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: RNA (genomic)
San Francisco
                               California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        linear
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                                  Sequence 0.0, Application US/08838545

Sequence No. 6046307

GENERAL INFORMATION:
APPLICANT: Wright, Woodring E.
APPLICANT: Orey, David R.
APPLICANT: Patery W.
APPLICANT: Patery W.
APPLICANT: Orey, David R.
APPLICANT: No. 6046307con, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embardance and Townsend and Crew LLP
STREET: Two Embardance and Townsend and Crew LLP
STREET: We Embarded and Townsend and Crew LLP
STREET: USA
COUNTRY: USA
COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPREMATING SYSTEM: PC-DOS/MS-DOS
SOFTWARR: Parantin Release #1 0 Naceic. #1 200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 40, Application US/09349532

Sequence 40, Application US/09349532

GENERAL INFORMATION:
APPLICANT: Shay, Jerry W.
APPLICANT: Shay, Jerry W.
APPLICANT: Corey, Micczyslaw A.
APPLICANT: Corey, David R.
APPLICANT: Corey, David R.
APPLICANT: No. 6294650ton, James C.
TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 60
CORRESPONDENCE ADDRESS: 7 Ownsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1; Length 25;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OPERATING SYSTEM: PC-DOS/MS-DUS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/838,545
FILING DATE: 09-APR-1997
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 31,944
REFERENCE/DOCKET NUMBER: 015389-001610US
TELEPHONE: (415) 576-0300
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 40:
SEQUENCE CHARACTERISTICS:
LENGTH: 25 base pairs
TYPE: nucleic acid
STRANDEDNESS: aingle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          7; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear
MOLECULE TYPE: RNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 72.0%
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-08-838-545-40
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                                                                                                                                                         DB 1; Length 28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/057,351
FILING DATE: 08-APR-1994
CLASSIFICATION NUMBER: US/09/057,351
FILING DATE: 08-APR-1994
PRIOR APPLICATION NUMBER: US/09/057,102
FILING DATE: 07-JUL-1994
PRIOR APPLICATION NUMBER: US/08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION NUMBER: US/08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION NUMBER: US/08/330,123
FILING DATE: 07-JUN-1995
ATTONNEY/AGENT INFORMATION:
AND ARE OF TAKEN 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Enbarcadero Center, Eighth Floor CITY: San Francisco STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 29, Application US/09057351
Patent No. 6548298
GENERAL INFORMATION:
APPLICANT: Ford, Juni
APPLICANT: Ford, Juni
APPLICANT: Ford, Walter
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              015389-000821US
                                                                                                                                                     Query Match 5.5%; Score 25; DB Best Local Similarity 100.0%; Pred. No. 52; Matches 25; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                             81 TTTGCTCCCCGCGCGCTGTTTTCT 105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 81 TTTGCTCCCCCCCCCCCTGTTTTCT 105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
WEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC_DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFRENCE/DOCKET NUMBER: 01536
TELECOMMUNICATION INFORMATION:
TELEPAK: (415) 576-0200
INPORMATION FOR SEQ ID NO: 29:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 100.
Matches 25; Conservative
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TYPE: nucleic acid
STRANDEDNESS: single
              TOPOLOGY: linear
                                                 MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-09-057-351-29/c
                                                                              US-08-472-802C-29
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COMPUTER: IBM PC Compatible
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC Compatible
COMPUTER: Patentin Release #1.0, Version #1.30
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/472,802C
FILING DATE: 07-JUL-1995
CLASSIFICATION NUMBER: US/08/272,102
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/330,123
FILING DATE: 07-JUL-1994
ATTORNEY/AGENT INFORMATION:
NAMME: SMITCH, William M.
REGISTRATION NUMBER: 15389-000820
TELECOMMUNICATION NUMBER: 15389-000820
TELECOMMUNICATION NUMBER: 15389-000820
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INCORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Villeponteau, Bryant
APPLICANT: Peng, Junii
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
FILING DATE: 07-JUL-1994

PRIOR APPLICATION DATA:
APPLICATION UNDRER:
FILING DATE: 27-OCT-1994

ATTORNEY/AGENT INFORMATION:
NAME: SECOFILA, JOHN R.
REGISTRATION NUMBER: 015389-000830US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0300
TELEPAX: (415) 676-0300
TELENGHH: 28 base pairs
LENGHH: 28 base pairs
TYBE: nucleic acid
STRANDEDNESS: single
TOPOLLOGY: linear
MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5.5%; Score 25; DB 100.0%; Pred. No. 52; ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            81 TTTGCTCCCCGCGCGCTGTTTTCT 105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TTTGCTCCCCGCGCGCTTTTTCT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; Sequence 29, Application US/08472802C; Patent No. 5958680
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 29
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 100.(
Matches 25; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 28 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   USA
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US-08-472-802C-29/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-08-482-115B-28
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APPLICANT: Shay, Jerry W.
APPLICANT: Wright, Woodring E.
APPLICANT: Wright, Woodring E.
APPLICANT: Platyszek, Mieczyslaw A.
APPLICANT: Orey, David Mieczyslaw A.
APPLICANT: No. 6294650ton, James C.
APPLICANT: No. 6294650ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by
TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 60
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5.3%; Score 24; DB 1; Length 24;
100.0%; Pred. No. 52;
tive 0; Mismatches 0; Indels
                                COMPUTER FEADABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/838,545
FILING DATE: 09-APR-1997
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/630,019
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INFORMATION:
NAME: SLOCATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 015389-001610US
TIELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/349,532
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: Townsend and Townsend and Crew LLI
STREET: Two Embarcadero Center, Bighth Floor
TTTE: San Prancisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            41 TTTGTCTAACCCTAACTGAGAAGG 64
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      other nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 25, Application US/09349532
Patent No. 6294650
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
INPORMATION FOR SEQ ID NO: 25:
SEQUENCE CHARACTERISTICS:
LENGTH: 24 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity 100.
Matches 24; Conservative
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DESCRIPTION:
DESCRIPTION:
DESCRIPTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY:
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COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
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                                                               Sequence 2, Application US/08770565
Patent No. 5846723
GENERAL INFORMATION:
APPLICANT: Kim, Nam Woo
APPLICANT: Kealey, James T.
APPLICANT: Perizan, Roaley
APPLICANT: Perizan, Roaley
TITLE OF INVENTION: Methods for Detecting the RNA Component of
TITLE OF INVENTION: Telomerase
NUMBER OF SEGURCES: 26
CORRESPONDENCE ADDRESS:
ADDRESSEE: TOWNSEND and TOWNSEND and CREW LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
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Patent No. 6046307
GENERAL INFORMATION:
APPLICANT: Wright, Woodring E.
APPLICANT: Piatyszek, Mieczyslaw A.
APPLICANT: Corey, David R.
APPLICANT: No.69, David R.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Peptide Nucleic Acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPTRY: USA

ZIP: 94111-3834

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
SUSTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/770,565
FILING DATE: 20-DEC.1996
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Stochella, John R.
REGISTRATION NUMBER: 015389-002300US
TELEBROCK/DOCKET NUMBER: 015389-002300US
TELEBROWNICATION INFORMATION:
TELEBROWNICATION INFORMATION:
TELEBRANE: 415-576-0300
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 30 base pairs
TYPE: mucleic acid
TYPE: mucleic acid
TYPE: mucleic acid
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STEET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                     STREET: Two Embarcadero Center, 8th Floor
CITY: San Francisco
STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            5.4%; Score 24.4; D
96.2%; Pred. No. 62;
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            355 CCTTTCAGGCCGCAGGAAGAGGAACG 380
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 96.2
Matches 25; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bingle
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; MOLECULE TYPE: DNA
US-08-770-565-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STRANDEDNESS:
                      RESULT 79
US-08-770-565-2/c
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                                                                                                                                                                                                                                                                                                                                                                                  STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ò
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/desc = "peptide nucleic acid (PNA), where (deoxy(ribose-phosphate linkages are replaced by N-(2-aminoethyl)glycine units linked to nucleotide bases via glycine amino N through a methylenecarbonyl linker" ö Gaps ö

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APPLICANT: Silverman, Robert H.

APPLICANT: Silverman, Robert H.

APPLICANT: Silverman, Robert H.

APPLICANT: Cowell, John K.

APPLICANT: Cowell, John K.

APPLICANT: Cowell, John K.

APPLICANT: Torrence, Paul F.

ITILE OF INVENTION: RNASE L ACTIVATORS AND ANTISENSE OLIGONUCLEOTIDES

ITILE OF INVENTION: EFFECTIVE TO TREAT TELOMERASE-EXPRESSING MALIGNANCIES

TITLE OF INVENTION: BFFECTIVE TO TREAT TELOMERASE-EXPRESSING MALIGNANCIES

CURRENT APPLICATION NUMBER: US/09/018,125A

CURRENT FILING DATE: 1999-02-03

EARLIER APPLICATION NUMBER: 60/044,507

EARLIER FILING DATE: 1997-04-21

NUMBER OF SEQ ID NOS: 9

SOFTWARE: PatentIN Ver. 2.0

LENGTH: 24
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JUS-08-838-545-26

JUS-08-838-545-26

JECTOR O. 604630TO:

JENERAL INPORMATION:

APPLICANT: Whight, Woodring E.

APPLICANT: Piatysek, Mieczyslaw A.

APPLICANT: Piatysek, Mieczyslaw A.

APPLICANT: Piatysek, Mieczyslaw A.

APPLICANT: Power, David R.

APPLICANT: Power, David R.

TITLE OF INVENTION: Modulation of Mammalian Telomerase by

TITLE OF INVENTION: Peptide Nucleic Acids

NUMBER OF SEQUENCES: 60

CORRESPONDENCE ADDRESS:

ADDRESSEE: Townsend and Townsend and Crew LLP

STREET: Two Embarcadero Center, Eighth Floor

CITY: San Francisco

CONPUTRY: USA

COMPUTRY: IBM PC Compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARRE: PACHALIN BATA:

SOFTWARRE: PACHALIN BATA:

PILLIANTON NUMBER: US/08/838,545

FILLIANG DATE:

APPLICATION NUMBER: US/08-ABS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:

COTHER INFORMATION: Description of Artificial Sequence: primer US-09-018-125-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 5.3%; Score 24; DB 1; Length 24; Best Local Similarity 100.0%; Pred. No. 52; Matches 24; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             015389-001610US
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FILING DATE: 09-APR-137,
CLASSIPICATION: 536
FRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-196
ATTORNY AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 015389-00:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA
ORGANISM: Artificial Sequence
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INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 84
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                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid STRANDEDNESS: single STRANDEDNESS: single STRANDEDNESS: single TOPOLOGY: linear TOPOLOGY: linear DESCRIPTION: desc = "peptide nucleic acid (PNA), DESCRIPTION: where (deoxy(ribose-phosphate linhages are replaced by DESCRIPTION: N-(2-aminoethy) glycine units linhed to nucleotide bases via DESCRIPTION: glycine amino N through a methylenecarbonyl linher"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Silverman, Robert H.
APPLICANT: Sowell, John K.
APPLICANT: Cowell, John K.
APPLICANT: Cowell, John K.
APPLICANT: Cowell, John K.
APPLICANT: Li, Guiying
APPLICANT: Torrence, Paul F.
ITILE OF INVENTION: RNASE L ACTIVATORS AND ANTISENSE OLIGONUCLEOTIDES
ITILE OF INVENTION: RNASE L ACTIVATORS AND ANTISENSE OLIGONUCLEOTIDES
ITILE OF INVENTION: RPFECTIVE TO TREAT TELOMERASE-EXPRESSING MALIGNANCIES
FILE REFERENCE: 8656-022
CURRENT APPLICATION NUMBER: US/09/018,125A
CURRENT APPLICATION NUMBER: 60/044,507
EARLIER FILING DATE: 1997-04-21
MUMBER OF SEQ ID NOS: 9
SOFTWARE: PATENTIN VOIL 2.0
SEQ ID NO 4
LENGTH: 24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 5.3%; Score 24; DB 1; Length 24; Best Local Similarity 100.0%; Pred. No. 52; Matches 24; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5.3%; Score 24; DB 1; Length 24;
100.0%; Pred. No. 52;
cive 0; Mismatches 0; Indels
  FILING DATE: 09-APR-1997

FILING DATE: 09-APR-1997

APPLICATION NUMBER: US 08/630,019

FILING DATE: 09-APR-1996

ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944

REFERENCE/DOCKET NUMBER: 015389-001610US

TELEPHONE: (415) 576-0200

TELEPAX: (415) 576-0200

TELEPAX: (415) 576-0300

INFORMATION FOR SEQ ID NO: 25:
SEQUENCE CHARACTERISTICS:
LENGTH: 24 base pairs
TYPE: nucleic acid
STRANDENNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  41 TTTGTCTAACCCTAACTGAGAAGG 64
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 83
US-09-018-125-5/c
; Sequence 5, Application US/09018125A
; Patent No. 6468983
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity 100.
Matches 24; Conservative
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-349-532-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-09-018-125-4
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US-08-485-778-20
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US-08-485-778-20
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                                                                                                                                                               bases via
                                                                                                                                                                                                                                                                                                         ö
                                                             TOPOLOGY: linear MOLECULE TYPE: other nucleic acid (PNA), DESCRIPTION: /desc = "peptide nucleic acid (PNA), DESCRIPTION: where (deoxy/ribose-phosphate linkages are replaced by DESCRIPTION: N-(2-aminoethyllglycine units linked to nucleotide base DESCRIPTION: glycine amino N through a methylenecarbonyl linker"
                                                                                                                                                                                                                                                                                                         Gaps
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APPLICANT: Wright, Woodring E.
APPLICANT: Piatyszek, Mieczyslaw A.
APPLICANT: Correy, David R.
APPLICANT: No. 6294650ton, James C.
ANDERSONO PORTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 60
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                      5.1%; Score 23; DB 1; Length 23; 100.0%; Pred. No. 58; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/349,532
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: Townsend and Townsend and Crew LLP STREEF: Two Enharcadero Center, Eighth Floor CITY: San Francisco STATE: California
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PRIOR APPLICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/838,545
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REGISTRATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFAN: (415) 576-0200
INFORMATION FOR SEQ ID NO: 26:
SEGUENCE CHARACTERISTICS:
LENGTH: 23 base pairs
TTYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                        35 CCATTTTTGTCTAACCTAACT 57
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Patent No. 6294650
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                            23; Conservative
23 base pairs
                      TYPE: nucleic acid
STRANDEDNESS: single
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DESCRIPTION:
DESCRIPTION:
DESCRIPTION:
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DESCRIPTION:
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                                                                                                                                                                                                            US-08-838-545-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-09-349-532-26
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                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Greider, Carol
APPLICANT: Marhuenda, Maria Antonia Blasco
APPLICANT: Marhuenda, Maria Antonia Blasco
APPLICANT: Villegonteau, Bryant
TITLE OF INVENTION: RNA COMPONENT OF TELOWERASE
NUMBER OF SEQUENCES: 45
CORRESPONDENCE ADDRESS:
ADDRESSEE: Two Allitia Drive
CITY: Lexington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION WHERE: US/08/485,778
FILING DATE: 07-JE-1995
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                              APPLICANT: Andrews, William H.
APPLICANT: Avilion, Ariel Athena
APPLICANT: Feng, Junli
APPLICANT: Funk, Walter
APPLICANT: Greider, Carol
APPLICANT: Marhuenda, Maria Antonia Blasco
Query Match 5.1%; Score 23; DB Best Local Similarity 100.0%; Pred. No. S8; Matches 23; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 5.1%; Score 23; DB Best Local Similarity 100.0%; Pred. No. 70; Matches 23; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION 1435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/387,524
FILING DATE: 13-FEB-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 07-OCT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
ATTORNEY/AGENT INFORMATION:
NAME: Granahan, Patricia
REGISTRATION NUMBER: 32,227
FELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFRAX: 617-861-6340
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         35 CCATTTTTGTCTAACCCTAACT 57
                                                                                                                            CCATTTTTGTCTAACCCTAACT 23
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                                                                                                                                                                                                                                                            ; Sequence 20, Application US/08485778
; Patent No. 5876979
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 02173
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: 617-861-9540
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 27 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
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4.9%; Score 22; DB 1; Length 22; 100.0%; Pred. No. 66; 0; Indels tive 0; Mismatches 0; Indels
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APPLICANT: FENG, Juni,
APPLICANT: FENG, Juni,
APPLICANT: FENG, Juni,
APPLICANT: FUNK, Walter
APPLICANT: FUNK, Walter
APPLICANT: FUNKWING HUMAN TELOMERASE
TITLE OF INVENTION: HUMAN TELOMERASE
CORRESPONDENCE: 25
CORRESPONDENCE: 25
CORRESPONDENCE: 15
ADDRESSE: Townsend and Townsend Khourie and Crew
STREET: 379 Lytton Avenue
CITY: Palo Alto
CONTENT: California
                                                                                                                                                                                                         COUNTER: US

ZITS 94301

COMPUTER READABLE FORM:
MEDIUM TYDE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/330,123A
FILING DATE: 27-0CT-1994
CLASSIFICATION NUMBER: US 08/272,102
FILING DATE: 07-UUL-1994
ATTORNEY/AGENT INPORMATION:
PROTESTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 15389-000810
TELEPONE: (415) 326-2420
TELEPONE: (415) 326-3420
                               CORRESPONDENCE ADDRESS:
ADDRESSE: Townsend and Townsend Khourie and Crew STREET: 379 Lytton Avenue CITY: Palo Alto STATE: California COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: BATENIN PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATE:
FILING DATE: 27-OCT-1994
FLING DATE: 77-OCT-1994
PRIOR APPLICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        46 CTAACCCTAACTGAGAAGGGCG 67
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; Sequence 6, Application US/08330123A
; Patent No. 5583016
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity 100. Matches 22; Conservative
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-08-330-123A-5
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                                                                                               PARENT NO. 601.148

APPLICANT: Andrews, William H.
APPLICANT: Avilion, Ariel A.
APPLICANT: Feng, Unni
APPLICANT: Feng, Unni
APPLICANT: Feng, Unni
APPLICANT: Feng, Unni
APPLICANT: Greider, Carol
APPLICANT: Greider, Carol
APPLICANT: Greider, Carol
APPLICANT: Greider, Carol
APPLICANT: Willeponteau, Bryant
TITLE OF INVENTION: RNA Component of Telomerase
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C.
STREET: Two Militia Drive
CITY: Lexington
STREET: Two Militia Drive
CITY: Lexington
STREET: RNA MILIA DRIVE
COMPUTER IN PROMBER: US/08/520,550A
COUNTRY: US
COMPUTER: IS PLOPPY disk
COMPUTER: IS PLOPPY disk
COMPUTER: LEMP PC COMPATIBLE
COMPUTER: LEMP PC COMPATIBLE
COMPUTER: SPEADABLE FORM:
MEDIUM TYPE: CP.055/MS-DOS
CNETWARE: 29-ANG-1995
FILING DATE: 13-FEB-1995
FILING 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  5.1%; Score 23; DB 1; Length 27; ilarity 100.0%; Pred. No. 70; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Indels
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Sequence 5, Application US/08330123A
Patent No. 5833016
GENERAL INFORMATION:
APPLICANT: VILLEPONTEAU, Bryant
APPLICANT: FENG, Junis
APPLICANT: FUNK, Walter
APPLICANT: FUNK, Walter
TITLE OF INVENTION: HUMAN TELOMERASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        22 GGAGGGTGGTGGCCATTTTTG 44
                                                 Sequence 20, Application US/08520550A
Patent No. 6013468
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tes 23; Conserva
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                 US-08-520-550A-20
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100.08;
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TELEPAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 27:
SEQUENCE CHARACTERISTICS:
                                                                                                                                   Query Match
Best Local Similarity 100.0
Matches 22; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 100.0
Marches 22; Conservative
                                                                                                                                                                                                                                                                      183 CTGCTGGCCCGTTCGC
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                linear
          TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 92
US-08-660-678A-28/c
                                                                          US-08-482-115B-32
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 91
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Sequence 32, Application US/08482115B
Patent No. 576679
GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Juni;
TITLE OF INVENTION: Assays for the RNA Component of Human
TITLE OF INVENTION: Assays for the RNA Component of Human
TITLE OF INVENTION: Telomerase
NUMBER OF SEQUENCES: 40
CORRESPONDENCE ADDRESS: 40
STREET: Townsend and Townsend and Crew LLP
STREET: Townsend and Townsend and Crew LLP
STREET: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 22; DB 1; Length 22;
Pred. No. 66;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: CALITOLILIA
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN DATA:
APPLICATION NUMBER: US/08/482,115B
FILING DATE: 07-UN-1995
CLASSIFTCATION 0ATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-UL-1994
FILING DATE: 07-UL-1995
FILING DATE: 07-UL-1994
FILING DATE: 07-UL-1994
FILING DATE: 07-UL-1994
FILING DATE: 07-UL-1994
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FILING DATE: 07-UL-1996
FILING DATE: 07-UL-1996
FILING DATE: 07-UL-1996
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100.0%; Pred. No. c...
0; Mismatches
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
ATYONEY/AGENT INFORMATION:
NAME: Smith, William M
REGISTRATION NUMBER: 30,233
REFERENCE/DOCKET NUMBER: 15389-000810
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFAX: (415) 326-2400
TELEFAX: (415) 326-3400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   54 AACTGAGAAGGGCGTAGGCGCC 75
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     22 AACTGAGAAGGGCGTAGGCGCC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 100.0
Matches 22; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             single
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STRANDEDNESS: sing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-08-482-115B-32/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-08-330-123A-6
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100.0%; Pred. No. 66;
DB 1; Length 22;
66;
                                                 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: CALLLIANTS

ZIP: 94111-3834

COUNTRY: USA

ZIP: 94111-3834

COMPUTER READABLE FORM:

MEDIUW TYPE: FORD

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/660,678A

FILING DATE: 05-JUN-1996

CLASSIFICATION: 435

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/330,123

FILING DATE: 27-OCT-1994

PRIOR APPLICATION NUMBER: US 08/320,123

FILING DATE: 07-JUL-1994

ATTORNEY/AGENT INFORMATION:

NAME: SLOCABILA, JOHN R.

REGISTRATION NUMBER: 32,944

REFERENCE/DOCKET NUMBER: 015389-000811US

TELECOMMUNICATION INFORMATION:

MATERIALION NUMBER: 32,944

REFERENCE/DOCKET NUMBER: 015389-000811US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco Center, Eighth Floor STATE: California
  4.9%; Score 22; DB 100.0%; Pred. No. 66; ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Feng, Junii
APPLICANT: Feng, Junii
APPLICANT: Funk, Walter
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Cr
                                                                                                CCCTCC 204
                                                                                                                                                                                                                               Sequence 27, Application US/08660678A Sequence 27, Application US/08660678A Setent No. 5837857 GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ... 46 CTAACCTAACTGAGAAGGGCG 67
                                                                                                                              22 CTGCTGGCCCGTTCGCCCCTCC 1
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Gaps
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                                                                                                                                 COMPUTER: FIDDY GISK
COMPUTER: FIDDY GISK
COMPUTER: IBM PC compatible
COMPUTER: BEACHLIN Release #1.0, Version #1.30
SOUTHARE: Patentin Release #1.0, Version #1.30
CUBRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/485,778
FILING DATE: 07-01-195
CHASSIFTCATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/387,524
FILING DATE: 13-FEB-195
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 07-01-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/72,102
FILING DATE: US 08/72,102
FILING DATE: US 08/72,102
RATCRINGY/AGENT INFORMATION:
NAME: Granahan, Particia
REGISTRATION NUMBER: 32,227
REFERENCE/DOCKET NUMBER: CSHL94-05A4
TELEPPAK: 617-861-6240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C. STREET: Two Militia Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Andrews, William H.
APPLICANT: Andrews, William H.
APPLICANT: Feng, Juni
APPLICANT: Feng, Juni
APPLICANT: Feng, Juni
APPLICANT: Greider, Carol
APPLICANT: Greider, Carol
APPLICANT: Marhuenda, Marria Antonia Blasco
APPLICANT: Willeponteau, Bryant
TITLE OF INVENTION: RNA COMPONENT OF TELOMERASE
NUMBER OF SEQUENCES: 45
CORRESPONDENCE ADDRESS:
ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.(
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATURE SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               46 CTAACCCTAACTGAGAAGGGCG 67
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: 7: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: single
                                                                         ZIP: 02173
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; TOPOLOGY: linear
US-08-485-778-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
Matches 22; Conserv
       Lexington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               02173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: Les
STATE: MA
COUNTRY:
                                                        COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 94
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       g
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Pred. No. 66;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Indels
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Juni
APPLICANT: Funk, Walter
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 30
CORRESPONDENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Bighth Floor
CITY: San Francisco
STATE: California
COUNTRY: USA
ZIP: 9411.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hamilton, Brook, Smith & Reynolds, P.C.
Two Militia Drive
                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATURG SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTION DATA:
APPLICATION NUMBER: US/08/660,678A
FILING DATE: 05-UN-1996
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Feng, Junia
APPLICANT: Funk, Walter
APPLICANT: Grader, Carol
APPLICANT: Grader, Carol
APPLICANT: Villeponteau, Bryant
TITLE OF INVENTION: RNA COMPONENT OF TELOWERASE
TOWNER OF SEQUENCES: 45
CORRESSPONDENCE ADDRESS:
ADDRESSEE: Hamilton, Brook, Smith & Reynolds,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
ATTORNEY AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944
RECISTRATION NUMBER: 32,944
TELEPHONE: (415) 576-0200
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 28:
SEQUENCE CHARACTERISTICS:
LENGTH : 22 base pairs
LENGTH : 22 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Andrews, William H.
Avilion, Ariel Athena
Feng, Junli
Funk, Walter
Greider, Carol
Marhuenda, Maria Antonia Blasco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4.9%; Score ... 66;
100.0%; Pred. No. 66;
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       Sequence 28, Application US/086606787
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AACTGAGAAGGGCGTAGGCGCC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 7, Application US/08485778; Patent No. 5876979; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 100.u
....hes 22; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
STRANDEDNESS: sing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear
MOLECULE TYPE: DNA
US-08-660-678A-28
                       Patent No. 5837857
GENERAL INFORMATION:
APPLICANT: Villepo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 93
US-08-485-778-7/c
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APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET:
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66;
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100.0%; Pred. No. 66;
cive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                               0; Indels
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CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/472,802C
FILING DATE: 07-JUN-1995
CLASSIETCATION: 514
PRIOR APPLICATION NUMBER: US/08/72,102
PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/330,123
FILING DATE: 07-OCT-1994
ATTORNEY/AGENT INFORMATION:

NAME: SMALM: William M.

REGISTRATION NUMBER: 15389-000820
TELECOMMUNICATION NUMBER: 15389-000820
TELECOMMUNICATION NUMBER: 15389-000820
TELECOMMUNICATION NUMBER: 15389-000820
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 42:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE TOWNERD OF SEQUENCE SEQUENCE 42, Application US/08472802C

Sequence 42, Application US/08472802C

Sequence 42, Application US/08472802C

RENEAL INFORMATION:

APPLICANT: Villeponteau, Bryant
APPLICANT: Villeponteau, Bryant
APPLICANT: Andrews, William H.

TITLE OF INVENTION: Mammalian Telomerase

NUMBER OF SEQUENCES: 44

CORRESPONDENCE ADDRESS:

ADDRESSEE: Townsend and Townsend and Crew LLP

STREET: Two Embarcadero Center, Eighth Floor

CITY: San Francisco

STATE: California
                                                                                                                                                                                                                                                                        4.9%; Score 22; DB 100.0%; Pred. No. 66; tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                               183 CTGCTGGCCCGTTCGCCCTCC 204
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          46 CTAACCCTAACTGAGAAGGGCG 67
                                                                                                                                                                                                                                                                                                                                                                                                                                crácreacecerreaceceree 1
                    TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 37
SEQUENCE CHARACTERISTICS:
LENGTH: 22 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 100.0
Marches 22; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 100.0
Matches 22; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 22 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: nucleic acid
STRANDEDNESS: sing
                                                                                                                                                                          TOPOLOGY: linear;
MOLECULE TYPE: DNA
US-08-472-802C-37
  TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                 à
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100.0%; Pred. No. 66;
iive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Riopy disk
COMPUTER: IBM PC compatible
SOFTWARE: Patentin Release #1.0, Version #1.30
CURENT APPLICATION DATA:
APPLICATION NUMBER: US/08/472,802C
FILING DATE: 07-JUN-1995
RILING DATE: 07-JUL-1994
PRIOR APPLICATION NUMBER: US 08/272,102
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
ATTORNEY/AGENT INFORMATION:
NAME: SMITH, William M.
REGISTRATION NUMBER: 30,223
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-08-472-802C-37/C

Sequence 37, Application US/08472802C

Sequence 37, Application US/08472802C

Sequence 37, Application US/08472802C

SEMERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 44

CORRESPONDENCE ADDRESSS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STRATE: California
COUNTRY: USA
                    FILING DATE: 07-0E-1995
CLASSIFICATION 1435
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
FILING DATE: 13-FEB-1995
FILING DATE: 13-FEB-1995
FILING DATE: 27-0CT-1994
PRIOR APPLICATION NUMBER: US 08/330,123
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-0CL-1994
ATTORNEY/AGENT INFORMATION:
NAME: Granahan, Parricia
REGISTRATION NUMBER: 32,227
REFERENCE/DOCKET NUMBER: CSHL94-05A4
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFRANCE: 617-861-6240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER: 15389-000820
FELECOMMUNICATION INFORMATION:
UMBER: US/08/485,778
07-JE-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        54 AACTGAGAAGGGCGTAGGCGCC 75
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AACTGAGAAGGGCGTAGGCGCC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO: 8: SEQUENCE CHARACTERISTICS: LENGTH: 22 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity 100.
Matches 22; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
STRANDEDNESS: single
  APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-08-485-778-8
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4.9%; Score 22; DB 1; Length 22;
100.0%; Pred. No. 66;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBW PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Petentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/520,550A
FILING DATE: 29-AuG-1995
CLASSIFTCATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 13-FEB-1995
FILING DATE: 27-OCT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 32,227
FILING DATE: 07-UUL-1994
ATTORNEY/AGENT INFORMATION:
NUMBE: GRANABAN: 32,227
REPERENCE/DOCKET NUMBER: 32,227
                           NUMBER OF SEQUENCES: 47
CORRESPONDENCE ADDRESS:
ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C.
STREET: Two Militia Drive
CITY: Lexington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C. STREET: Two Militia Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Andrews, William H.
APPLICANT: Avilion, Ariel A.
APPLICANT: Feng, Junli
APPLICANT: Funk, Walter
APPLICANT: Funk, Walter
APPLICANT: Marhuenda, Maria A. B.
APPLICANT: Marhuenda, Maria A. B.
APPLICANT: Marhuenda, Maria A. B.
APPLICANT: Willeponceau, Bryant
TITLE OF INVENTON: RNA COMPONENT OF SEQUENCES: 47
CORRESPONDENCES ADRESS:
TITLE OF INVENTION: RNA Component of Telomerase NUMBER OF SEQUENCES: 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         46 CTAACCCTAACTGAGAAGGGCG 67
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; Sequence 8, Application US/08520550A
; Patent No. 6013468
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity 100.
nes 22; Conservative
                                                                                                                                                                                                COUNTRY: US
ZIP: 02173
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: nucleic acid
STRANDEDNESS: single
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COMPUTER READABLE FORM:
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/472,802C
FILING DATE: 07-UJN-1995
FILING PAPLICATION S14
PRIOR APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-UJL-1994
PRIOR APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M.
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 15389-000820
TELEFPAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 43:
SEQUENCE CHARACTERISTICS:
LENGTH: 22 base pairs
                                                                                                                           US-08-472-902C-43/C

1 Sequence 43, Application US/08472802C

1 Patent No. 5958680

1 GENERAL INFORMATION:

APPLICANT: Villeponteau, Bryant
APPLICANT: Peng, Junii
APPLICANT: Andrews, William H.

TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 44

CORRESPONDENCE ADDRESS:

ADDRESSE: Townsend and Townsend and Crew LLP

STRET: Two Embarcadero Center, Eighth Floor

CITY: San Francisco

CONTRIBER California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 98
US-08-520-550A-7/C
Sequence 7, Application US/08520550A
Patent No. 6013468
GENERAL INFORMATION:
APPLICANT: Andrews, William H.
APPLICANT: Avilion, Ariel A.
APPLICANT: Feng, Junil
APPLICANT: Feng, Junil
APPLICANT: Greider, Carol
APPLICANT: Marhuenda, Maria A. B.
APPLICANT: Walter
APPLICANT: Walter
APPLICANT: Villeponteau, Bryant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Andrews, William H.
Avillon, Ariel A.
Feng, Junii
Funk, Walter
Greider, Carol
Marhuenda, Maria
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              54 AACTGAGAAGGGCGTAGGCGCC 75
                  22 craacccraacrgagaagggg 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: nucleic acid STRANDEDNESS: sing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; MOLECULE TYPE: RNA
US-08-472-802C-43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: San Fra
STATE: Califo
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patemetr Release #1.0, Version #1.30
CURRATI APPLICATION DATA:
APPLICATION NUMBER: US/08/998,443
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco STATE: California
  PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-011-1994
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-000811US
TELEFORMUNICATION INFORMATION:
TELEFORMUNICATION INFORMATION:
TELEFORM (415) 576-0200
TELEFORM: (415) 576-0300
TELEFORM (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-08-1998-443-28/C
; Sequence 28, Application US/08998443
; Patent No. 6054575
; GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junii
APPLICANT: Funk, Walter
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: SCORELLA, JOHN R.
REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 28:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRICK APPLICATION DATA:
APPLICATION NUMBER: US/08/660,678
FILING DATE: 05-UN-1996
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRICK APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-ULL-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         46 CTAACCCTAACTGAGAAGGGCG 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CTAACCCTAACTGAGAAGGGCG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 100.
Matches 22; Conservative
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ADDRESSEE: Townsend
                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-998-443-27
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/520,550A
FILING DATE: 29-ANG-1995
CLASSIFICATION DATA:
APPLICATION NUMBER: US/08/320,524
FILING DATE: 11-FEB-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/387,524
FILING DATE: 12-CCT-1994
PRIOR APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
ATORNEY/AGENT INFORMATION:
NAME: GRANDANIND NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
ATORNEY/AGENT INFORMATION:
NAME: GRANDANIND NUMBER: 32,227
REFERENCE/DOCKET NUMBER: 32,227
REFERENCE/DOCKET NUMBER: CSHL94-05A3B
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 27, Application US/08998443

Sequence 27, Application US/08998443

GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junli
APPLICANT: Mammalian Telomerase
NUMBER OF SEQUENCES: 30
CORRESPONDENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
COUNTRY: USA
ZIP: 94111-3834
COMPUTER: READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARF: Defenter PC-DOS/MS-DOS
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APPLICATION NUMBER: US/08/998,443 FILING DATE:
CLASSIFICATION DATA:
APPLICATION NUMBER: US/08/660,678 FILING DATE: 05-JUN-1996 APPLICATION NUMBER: US/08/330,123 FILING DATE: 27-OCT-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    54 AACTGAGAAGGGCGTAGGCGCC 75
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: 617-861-9540
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
LENGTH: 22 base pairs
TYPE: nucleic acid
STRANDENNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 100.C
Matches 22; Conservative
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TOPOLOGY:
US-08-520-550A-8
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Sequence 6, Application US/09580517
                                      Patent No. 6320039
GENERAL INFORMATION:
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ZIP: 94301

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
SOFTWARE: PATENTIN ROLASE: US/09/580,517
FILING DATE: 25-M3y-2000
CLASSIFICATION NUMBER: US/09/580,517
FILING DATE: 400Mnown>
PRIOR FILING DATE: 400Mnown>
PRIOR FILING DATE: 400Mnown>
ATTORNEY/AGBNT INFORMATION:
NAME: Smith, William M
REGISTRATION NUMBER: 30,23
REFERENCE/DOCKET NUMBER: 15389-000810
TELEPAX: (415) 326-2422
; INFORMATION FROM INFORMATION:
TELEPAX: (415) 326-2422
; INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERIZES:
SEQUENCE: SEQUENCE: AND ASSECTION NO. 5:
SEQUENCE: SEQUENCE: ASSECTION NO. 5:
SEQUENCE: SEQUENCE: ASSECTION: ASSECTION NO. 5:
SEQUENCE: SEQUENCE: ASSECTION NO. 5:
SECUENCE: ASSECTION NO. 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend Khourie and Crew
                                                                                                                                                                                                                     4.9%; Score 22; DB 1; Length 22; 100.0%; Pred. No. 66; 0; Indels tive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ANDREWS, William H.
TITLE OF INVENTION: HUMAN TELOMERASE
NUMBER OF SEQUENCES: 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear

MOLECULE TYPE: RNA

SEQUENCE DESCRIPTION: SEQ ID NO: 5:
US-09-580-517-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: VILLEPONTEAU, Bryant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET: 379 Lytton Avenue
CITY: Palo Alto
STATE: California
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US-09-580-517-5/c
; Sequence 5, Application US/09580517
; Patent No. 6320039
; GENERAL INFORMATION:
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TYPE: nucleic acid
STRANDEDNESS: single
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FUNK, Walter
                                                                                                                                                                                                                                                        Best Local Similarity 100.0
Matches 22; Conservative
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Matches 22; Conservative
nucleic acid
                               STRANDEDNESS: sin
TOPOLOGY: linear
MOLECULE TYPE: DNA
                                                                                                                ; MOLECULE TY|
US-08-998-443-28
                                                                                                                                                                                                                         Query Match
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RESULT 103 US-09-580-517-6/c

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GENERAL INFORMATION:

APPLICANT LALEPOWIERJU Bryant
FREG, Junii,
NORMA, Maletrian H
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RESULT 107
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                                                                                                                                                                                       NAME/KEY: misc feature

LOCATION: (1) —

DOTHER INFORMATION: Biotin 5'-terminal

OTHER INFORMATION: Description of Artificial Sequence: Affinity Agent

US-09-717-8288-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: Biotin 5'-terminal
OTHER INFORMATION: Description of Artificial Sequence: Affinity Agent
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Weinrich, Scott L
APPLICANT: Atkinson III, Edward M
APPLICANT: Lichteteiner, Serge P
APPLICANT: Userserot, Allain P
APPLICANT: Vascorot, Ronald A
TITLE OF INVENTION: A Method for Purifying Telomerase
CURRENT APPLICATION NUMBER: US/09/17,829A
FILE REFERENCE: PurifiedTelomeraseOllbase
CURRENT APPLICATION NUMBER: 09/420,056
PRIOR APPLICATION NUMBER: 08/833,377
PRIOR APPLICATION NUMBER: 08/833,377
PRIOR APPLICATION NUMBER: 08/833,377
PRIOR FILING DATE: 1999-10-18
PRIOR FILING DATE: 1995-08-04
NUMBER OF SEQ ID NOS: 11
SOFTWARE: Patentin Ver. 2.1 edited
SEQ ID NO 2
MUMBER: ALENGTH: 22
                                                                                                                                                                                                                                                                                                                                                                                                                                  46 CTAACCCTAACTGAGAAGGGCG 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            46 CTAACCCTAACTGAGAAGGGCG 67
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; Sequence 2, Application US/09717829A
; Setten No. 6545133
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22 CTAACCCTAACTGAGAAGGGCG 1
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Patent No. 6548298
GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junii
APPLICANT: Feng, Junii
                      NUMBER OF SEQ ID NOS: 11
SOFTWARE: Patentin Ver. 2.1 edited
SEQ ID NO 2
LENGTH: 22
                                                                                                                       TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA ORGANISM: Artificial Sequence
PRIOR FILING DATE: 1995-08-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity 100.
Matches 22, Conservative
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LOCATION: (1)
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US-09-057-351-41/c
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US-09-717-829A-2
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                                                                                                                                                                        FEATURE:
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Gaps
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66;
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                                                                                                                                                                                                                                  CALF: JALE STATE COMPUTER REALABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/057,351
FILING DATE: 08-APR-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOWNERS:

STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
COUNTRY: USA
APPLICANT: Andrews, William H.

TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION 458
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-UUL-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION NUMBER: US 08/472,802
FILING DATE: 07-UN-1995
ATTORNEY/AGENT INFORMATION:
NAME: SECATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-000821US
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION SEQ ID NO: 41:
SEQUENCE CHARACTERISTICS:
LENGTH: 22 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junii
APPLICANT: Funk, Walter
APPLICANT: Andrews, Walliam H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            67
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; Sequence 42, Application US/09057351
Sequence 10, Application US/09057351
Patent No. 6548298
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         46 CTAACCCTAACTGAGAAGGGCG
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Best Local Similarity 100.0%; P
Matches 22; Conservative 0;
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear MOLECULE TYPE: RNA
                                                                                                                                                                                                            ZIP: 94111-3834
                                                                                                                                                                                                     USA
                                                                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-09-057-351-41
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100.0%; Pred. No. 74;
4.9%; Score 22; DB 1; Length 22; 100.0%; Pred. No. 66; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPRESS READABLE FORM:

COMPUTER READABLE FORM:

MEDLIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
CONFERNT APPLICATION DATA:
APPLICATION NUMBER: US/08/330,123A
FILING DATE: 27-0CT-1994
FILING DATE: 27-0CT-1994
FILING DATE: 37-0CT-1994
ATTORNEY/AGENT INFORMATION:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-0UL-1994
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M
REGISTRATION NUMBER: 15389-000810
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION OF 25:
SEQUENCE CHRRACTERISTICS:
INFORMATION FOR SEQ ID NO: 25:
SEQUENCE CHRRACTERISTICS:
LEMTH: 21 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: Townsend and Townsend Khourie and Crew STREET: 379 Lytton Avenue CITY: Palo Alto STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%; Prec. ....
                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: VILLEPONTEAU, Bryant
APPLICANT: FENG, Junil
APPLICANT: FUNK, Walter
APPLICANT: ANDREWS, William H.
TITLE OF INVENTION: HUMAN TELOMERASE
NUMBER OF SEQUENCES: 25
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend Khou
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 110
US-08-482-115B-25/C
Sequence 25, Application US/08482115B
Sequence 25, Application US/08482115B
Sequence 25, Application US/08482115B
GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Funk, Walter
APPLICANT: Funk, Walter
APPLICANT: Andrews, William H.
                                                                                                                                                                                                                                                                                                                                                 US-08-330-123A-25/c; Sequence 25, Application US/08330123A; Patent No. 5583016
                                                                                                                                              46 CTAACCCTAACTGAGAAGGGCG 67
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Best Local Similarity 100.(
Matches 21; Conservative
                                                                          22; Conservative
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EDNESS: single
       Query Match
Best Local Similarity
Matches 22; Conserva
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY:
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GENERAL INFORMATION:

APPLICANT: Geron Corporation
APPLICANT: Acting Scott
APPLICANT: Acting III, Edward
APPLICANT: Lichteteiner, Serge
APPLICANT: Lichteteiner, Serge
APPLICANT: Userson III, Edward
APPLICANT: Userson Ronald
TITLE OF INVENTION: Using Purified Telomerase to Identify Telomerase Activators and
TITLE OF INVENTION: Using Purified Telomerase to Identify Telomerase Activators and
TITLE OF INVENTION: Using Purified Telomerase to Identify Telomerase Activators and
TITLE OF INVENTION: Using Purified Telomerase to Identify Telomerase Activators and
TITLE OF INVENTION: Using Purified Telomerase to Identify Telomerase Activators and
TITLE OF INVENTION: Using Purified Telomerase to Identify Telomerase Activators and
TITLE OF INVENTION NUMBER: US/10/330,872
PRIOR FILING DATE: 1995-08-04
PRIOR FILING DATE: 1995-08-04
PRIOR FILING DATE: 1997-04-04
PRIOR FILING DATE: 2000-11-20
NUMBER OF SEQ ID NOS: 11
SOFTWARE: PatentIn version 3.1
SSPU ID NO 2
LENGTH: 22
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              COMPUTER: IBM FC COMPOSED

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Rolease #1.0, Version #1.30

SOFTWARE: Patentin Rolease #1.0, Version #1.30

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/057,351

FILING DATE: 04-APR-1994

PRIOR APPLICATION DATE: 07-JUL-1994

PRIOR APPLICATION DATE: 07-JUL-1994

PRIOR APPLICATION NUMBER: US 08/330,123

FILING DATE: 27-OCT-1994

PRIOR APPLICATION NUMBER: US 08/472,802

FILING DATE: 27-OCT-1994

PRIOR APPLICATION NUMBER: US 08/472,802

FILING DATE: 07-JUN-1995

APPLICATION NUMBER: US 08/472,802

FILING DATE: US US 08/472,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 100.0%; Pred. No. 00, Natches 22; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; Sequence 2, Application US/10330872; Patent No. 6787133
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CORGANISM: Homo sapiens
US-10-330-872-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear;
MOLECULE TYPE: RNA
US-09-057-351-42
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4.7%; Score 21; DB 1; Length 21; 100.0%; Pred. No. 74;
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; Sequence 33, Application US/08485778
; Patent No. 5876979
; GENERAL INFORMATION:
    APPLICANT: Avilion, Ariel Athena
    APPLICANT: Peng, Junil
    APPLICANT: Peng, Junil
    APPLICANT: Greider, Carol
    APPLICANT: Martuer
    APPLICANT: Greider, Carol
    APPLICANT: Martuer
    APPLICANT: Martuer

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MEDIUM TYPE: Floppy disk
COMPUTER: TBM PC compatible
OPERATUR SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
PFILING DATE: 07-JE-1995
CLASSIFICATION: 435
                PILING DATE: 05-JUN-1996

PILING DATE: 05-JUN-1996

CLASSIFTCATION NUMBER: US/08/660,678A

PILING DATE: 05-JUN-1996

PROR APPLICATION DATA:

APPLICATION NUMBER: US 08/330,123

PILING DATE: 07-JUL-1994

PRIOR APPLICATION NUMBER: US 08/272,102

PILING DATE: 07-JUL-1994

ATTORNEY AGENT INFORMATION:

NAME: SCOREILA, JOHN R.

REGISTRATION NUMBER: 32,944

REGISTRATION NUMBER: 32,944

REFERENCE/DOCKET NUMBER: 015389-000811US

TELEPHONE: (415) 576-0200

TELEPHONE: (415) 576-0300

INFORMATION FOR SEQ ID NO: 25:

SEQUENCE CHARACTERISTICS:

LENGTH: 21 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%; Prea. ...
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CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/387,524
FILING DATE: 13-FEB-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           184 TGCTGGCCCGTTCGCCCCTCC 204
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Best Local Similarity 100.0
Matches 21; Conservative
CURRENT APPLICATION DATA: APPLICATION NUMBER: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; TOPOLOGY: li;
; MOLECULE TYPE:
US-08-660-678A-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  S
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Assays for the RNA Component of Human Telomerase
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100.0%; Pred. No. 74;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                  COUNTRY: USA
ZIP: 9411-3834

ZIP: 9411-3834

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/482,115B
FILING DATE: 07-JUN-1995
FILING DATE: 07-JUN-1995
FILING DATE: 07-JUL-1994
FRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-0CT-1994
ATTORNEY/AGENT INFORMATION:
APPLICATION NUMBER: 32,944
FILING DATE: 27-0CT-1994
ATTORNEY/AGENT INFORMATION:
TELEPHONE: 27-0CT-1994
ATTORNEY/AGENT INFORMATION:
TELEPHONE: 415) 576-0200
INFORMATION FOR SEQ 100
INFORMATION FOR SEQ 100
INFORMATION FOR SEQ 100
INFORMATION CALION INFORMATION:
TELEPHONE: 415) 576-0200
INFORMATION FOR SEQ 100
INFORMATION FOR SEQ 100
INFORMATION CALION INFORMATION:
TYPE: ULBUSHESS: SINGLE
TYPE: TYPE: INFORMATION ATTORNEY
TYPE: TYPE: INFORMATION CALION C
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Patent No. 5837867
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Juni;
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PatentIn Release #1.0, Version #1.30
                                                                     NUMBER OF SEQUENCES: 40
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: San Francisco
STATE: California
COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, v
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           184 TGCTGGCCCGTTCGCCCCTCC 204
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Matches 21; Conservative
                                                                                                                                                                           STREET: TWO BURNESS CITY: San Francisco STATE: California
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 111
US-08-660-678A-25/c
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Gaps

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4.7%; Score 21; DB 1; Length 21; 100.0%; Pred. No. 74; iive 0; Mismatches 0; Indels

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Query Match
Best Local Similarity 100.C
Matches 21, Conservative
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CITY: Lexington
STATE: MA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; TOPOLOGY:
US-08-520-550A-33
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US-08-472-802C-26/C

# Sequence 26, Application US/08472802C

# Sequence 26, Application US/08472802C

# GENERAL INFORMATION:

# APPLICANT: Feng, Juni

# APPLICANT: Feng, Juni

# APPLICANT: Peng, Juni

# TITLE OF INVENTION: Mammalian Telomerase

# CORRESPONDENCE ADDRESS:

# ADDRESSEE: Townsend and Townsend and Crew LLP

# STREET: Two Embarcadero Center, Eighth Floor

# COUNTRY: San Francisco

# STATE: California

# COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:

POFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/472,802C
FILING DATE: 07-JUN-1995
CLASSIFICATION DATA:

APPLICATION DATA:

APPLICATION DATA:

PRIOR APPLICATION DATA:

APPLICATION DATA:

APPLICATION DATA:

APPLICATION NUMBER: US 08/372,102
FILING DATE: 07-JUL-1994

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994

ATTORNEY/AGENT INFORMATION:

NAME: SMICH William M.

REGISTRATION NUMBER: 15389-000820

TELEPRONE: (415) 576-0200

TELEPRONE: (415) 576-0300

INFORMATION FOR SEQ ID NO: 26:
SEQUENCE CHARACTERISTICS:
LENGTH: 21 base pairs

TYPE: MODILOR: 110-21
FILING DATE: 07-JUL-1994
ATTORNEY/AGENT INFORMATION:
NAME: Granahan, Patricia
REGISTRATION NUMBER: 32,227
REFERENCE/DOCKET NUMBER: CSHL94-05A4
TELECOMMUNICATION INFORMATION:
TELEPAK: 617-861-9540
INFORMATION FOR SEQ ID NO: 33:
SEQUENCE CHARACTERISTICS:
LENGTH: 21 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP. 34111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                              184 TGCTGGCCCGTTCGCCCTCC 204
                                                                                                                                                                                                                                                                                            ; TOPOLOGY: linear
US-08-485-778-33
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RESULT 115

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APPLICANT FENG, Junii
FUNK, Walter
ANDREWS, William H.
TITLE OF INVENTION: HUMAN TELOMERASE
NUMBER OS SEQUENCES: 25
CARESPONDENCE ADDRESS:
CARESPONDENCE TOwnsend and Townsend Khourie and Crew
STREET: 379 Lytton Avenue
CITY: Palo Alto
STATE: California
COMPUTRY: US
ZIP: 94.01
COMPUTRE READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER READABLE FORM:
COMPUTER: IBM PC Compatible
COMPUTER: PROPERING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PRECENTIN Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4.7%; Score 21; DB 1; Length 21;
100.0%; Pred. No. 74;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Indels
                                                                                       COMPUTER READABLE FORM:
MEDIUM TYBE: Ploppy disk
COMPUTER: IBM FC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/060,523
FILING DATE: 14-APR-1998
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/60,678
FILING DATE: 05-JUN-1996
FILING DATE: 27-OCT-1994
FILING DATE: 27-OCT-1994
FILING DATE: US 08/272,102
FILING DATE: US-1994
FILING DATE: US-1994
APPLICATION NUMBER: US 08/272,102
FILING DATE: US-1994
ATTORNEY/AGENT INFORMATION:
NAME: SCOTCHIA, JOHN R.
REGISTRATION NUMBER: 32,944
RESTERENCE/DOCKET NUMBER: 015389-000813US
TELEPHONE: (415) 576-0200
TELEPHONE: (415) 576-0200
TELEPAX: (415) 576-0200
TTYPE: MUCLEIC SEQIED
TYPE: MUCLEIC SEGIED
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iive 0; Mismatches
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APPLICATION NUMBER: US/09/580,517
FILING DATE: 25-May-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    184 TGCTGGCCCGTTCGCCCTCC 204
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; Sequence 25, Application US/09580517; Patent No. 6320039; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 100.
Matches 21; Conservative
                                                                   ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
San Francisco
California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  , MOLECULE TYPE: DNA
US-09-060-523-25
                                                  USA
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                                               COUNTRY:
ZIP: 941
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                                                                                       APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Juni
APPLICANT: Funk, Walter
APPLICANT: Funk, Walter
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                      COUNTY: USA
ZIP: 94111-3834
ZIP: 94111-3834
ZOMPUTER READABLE FORM:
MEDIUM TYEE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
ZURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/998,443
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  E: Townsend and Townsend and Crew LLP
Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AFPLICATION NUMBER: US/US/399,443
FILING DATE:
CLASSIFICATION DATA:
APPLICATION NUMBER: US/08/660,678
FILING DATE: 05-UN-1996
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-UL-1994
ATTONNEY AGENT INFORMATION:
NAME: SCOFELLA, JOHN R.
REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944
RECECOMMUNICATION INPORMATION:
TELEPHONE: (415) 576-0300
INPORMATION FOR SEQ ID NO: 25:
SEQUENCE CHARACTERISTICS:
LENGTH: 21 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 25, Application US/09060523
Sequence 25, Application US/09060523
GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Funk, Walter
TITLE OF INVENTION: Mammalian Telomerase
TITLE OF INVENTION: Mammalian Telomerase
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Crew
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 184 TGCTGGCCCGTTCGCCCTCC 204
US-08-998-443-25/c
; Sequence 25, Application US/08998443
; Patent No. 6054575
; GENERAL INFORMATION:
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STRANDEDNESS: single
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            , MOLECULE TYPE: DNA
US-08-998-443-25
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US-09-060-523-25/c
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OTHER INFORMATION: /mod base= OTHER OTHER INFORMATION: /note= "N = 5' biotinylated cytidine"
                                                                                                                                                                                  4.7%; Score 21; DB 1; Length 21; 100.0%; Pred. No. 74; cive 0; Mismatches 0; Indels
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Sequence 3, Application US/0883377

Patent No. 5968506

GENERAL INFORMATION:
APPLICANT: Weinrich, Scott L.
APPLICANT: Atkinson III, Edward M.
APPLICANT: Lichtsteiner, Serge P.
APPLICANT: Pruzan, Ronald A.
APPLICANT: Pruzan, Ronald A.
APPLICANT: Pruzan, Pruzan, Pruzan, Forlain P.
ITILE OF INVENTION: Purified Telomerase
NUMBER OF SEQUENCES: 15
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Crew LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: -
LOCATION: 1..22
OTHER INFORMATION: /note= "oligonucleotide P3"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: Two Embarcadero Center, Bighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                 184 TGCTGGCCCGTTCGCCCTCC 204
                                     LENGTH: 21 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLGY: linear
MOLECULE TYPE: DNA
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NAME/KEY: modified_base
                                                                                                                                                                                                     Best Local Similarity 100.
Matches 21; Conservative
  INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: DNA
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                                                                                                                                            US-09-057-351-25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: USA

ZIP: 94111-3834

COMPUTER READABLE FORM:
MEDIUW TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM:
PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/057,351
FILING DATE: 03-APR-1994
CLASSIFICATION: 435
PRIOR APPLICATION NUMBER: US/09/057,102
FILING DATE: 07-JUL-1994
PRIOR APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION NUMBER: US 08/310,123
FILING DATE: 07-JUL-1994
PRIOR APPLICATION NUMBER: US 08/310,123
FILING DATE: 107-JUN-1995
ATTORNEY/AGENT INFORMATION:
ANAMER APPLICATION LABORATION:
ANAMER APPLICATION NUMBER: US 08/472,802
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 118
US-09-057-351-25/c
; Sequence 25, Application US/09057351
; Patent No. 6548298
; GENERAL INFORMATION:
; APPLICANT: Villeponteau, Bryant
APPLICANT: Funk, Walter
; APPLICANT: Funk, Walter
; TITLE OF INVENTION: Mammalian Telomerase
; NUMBER OF SEQUENCES: 42
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
; CITY: San Francisco
        PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/330,123
FILING DATE: «Unknown»
ATTORNY/AGENT INFORMATION:

NAME: Smith, william M
REGISTRATION NUMBER: 30,23
REFERENCE/DOCKET NUMBER: 15389-000810
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 326-2400
TELEFAX: (415) 326-2422
INFORMATION FOR SEQ ID NO: 25:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE, DOCKET NUMBER: 015389-000821US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576.0200
TELEFAX: (415) 576.0200
                                                                                                                                                                                                                                                                                                   ; TOPOLOGY: linear
; MOLECULE TYPE: DNA
; SEQUENCE DESCRIPTION: SEQ ID NO: 25:
US-09-580-517-25
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        184 TGCTGGCCCGTTCGCCCTCC 204
                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 21 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                21; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity
Matches 21; Conserva
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US-08-482-115B-7/c
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                                                                                          Gaps
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                                             DB 1; Length 22; 78;
                                                                                          0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:

MEDIUM TYPE: S1097 disk

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: APELICATION DATA:

APPLICATION NUMBER: US/08/330,123A

FILING DATE: 27-CCT-1994

CLASSIFICATION: 435

PRIOR APPLICATION NUMBER: US 08/272,102

FILING DATE: US 08/272,102

ATTORNEY/AGENT INFORMATION:

NAME: SMIGHT MINIBLE: 30,223

REFERENCE/DOCKET NUMBER: 15389-000810

TELECOMMUNICATION INFORMATION:

MEDIUM TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:

THE TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                          APPLICANT: VILLEBONTEAU, Bryant
APPLICANT: FENG, Junii
APPLICANT: FUNK, Walter
APPLICANT: ANDREWS, William H.
TITLE OF INVENTION: HUMAN TELOMERASE
NUMBER OF SEQUENCES: 25
CORRESPONDENCE ADDRESS:
ADDRESSE: Townsend and Townsend Khourie and Crew
                                          4.7%; Score 21; DB 100.0%; Pred. No. 78; tive 0; Mismatches
                                                                                                                                                                                                                                                                                             Sequence 4, Application US/08330123A Patent No. 5583016 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-08-330-123A-7/c
; Sequence 7, Application US/08330123A
; Patent No. 5583016
; GENERAL INFORMATION:
APPLICANT: VILLEPONTEAU, Bryant
APPLICANT: FENG, Junil
                                                                                                                                  46 CTAACCCTAACTGAGAAGGGC 66
                                                                                                                                                               22 CTAACCCTAACTGAGAAGGGC 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 379 Lytton Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (415) 326-2400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: (415) 336-2422
INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                      Query Match
Best Local Similarity 100.0
Matches 21, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 100.
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: Palo Alto
STATE: California
COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear MOLECULE TYPE: RNA US-08-330-123A-4
                                                                                                                                                                                                                                                  RESULT 120
US-08-330-123A-4/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE:
US-08-833-377-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 8
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APPLICANT: Feng, Junia
APPLICANT: Feng, Junia
APPLICANT: Feng, Junia
APPLICANT: Andrews William H.
APPLICANT: Andrews William H.
TITLE OF INVENTION: Assays for the RNA Component of Human
TITLE OF INVENTION: Telomerase
NUMBER OF SEQUENCES: 40
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1; Length 20;
83;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Indels
                                                                                                                                                                                         APPLICANT: FUNK, Walter
APPLICANT: ANDREWS, William H.
TITLE OF INVENTION: HUMAN TELOMERASE
NUMBER OF SEQUENCES: 25
CORRESPONDENCES: 25
CORRESPONDENCES: 15
ADDRESSEE: Townsend and Townsend Khourie and Crew
STREET: 379 Lytton Avenue
CITY: Palo Alto
STATE: US
COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: ISM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4.4%; Score 20; DB 100.0%; Pred. No. 83; ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 7, Application US/08482115B
Patent No. 5776679
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2 GGTTGCGGAGGGTGGGCCTG 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: (415) 326-2400
TELEFAX: (415) 326-2422
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 100.0
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: USA
ZIP: 94111-3834
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Gaps
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83;
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                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Indels
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ZIP: 94111-3834

ZIP: 94111-3834

COMPUTER READALE FORM:

MEDIUW TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PATENTIN Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/660,678A

FILING DATE: 05-UN-1996

CLASSIFICATION 1435

PRIOR APPLICATION NUMBER: US/08/330,123

FILING DATE: 27-OCT-1994

PRIOR APPLICATION NUMBER: US/08/330,123

FILING DATE: 27-OCT-1994

PRIOR APPLICATION NUMBER: US/08/330,123

FILING DATE: 27-OCT-1994

ATPONENTY/AGENT INFORMATION:

ANAME: ALCONING ALCONING ALCONING ANAME: ACCOUNTRY/AGENT INFORMATION:

ACCOUNTRY/AGENT INFORMATION:

ANAME: ACCOUNTRY/AGENT INFORMATION:

ACCOUNTRY/AGENT INFORMATION:

ACCOUNTRY/AGENT INFORMATION:

ANAME: ACCOUNTRY/AGENT INFORMATION:

ANAME: ACCOUNTRY/AGENT INFORMATION:

ACCOUNTRY/AGENT 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          E: Townsend and Townsend and Crew LLP
Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                      Query Match

4.4%; Score 20; DB
Best Local Similarity 100.0%; Pred. No. 83;
Matches 20; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Juni,
APPLICANT: Funk, Walter
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-660-678A-26/c
; Sequence 26, Application US/08660678A
; Patent No. 5837857
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2 GGTTGCGGAGGGTGGGCCTG 21
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REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 01
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0200
INFORMATION FOR SEQ ID NO: 26: SEQUENCE CHARACTERISTICS:
   (415) 576-0300
                              INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 100.
Matches 20; Conservative
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                                                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: single
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California
                                                                                                                                                                                                                                                              , MOLECULE TYPE: DNA
US-08-660-678A-7
                                                                                                                                                                                                                               linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE:
                                                                                                                                                                                                                               TOPOLOGY:
   TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET:
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Pred. No. 83;
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/660,678A
FILING DATE: 05-JUN-1996
CLASSIFICATION: 435
PRIOR APPLICATION NUMBER: US/08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
ATORNEY/AGENT INPORMATION:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
ATORNEY/AGENT INPORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REBRERNCE/POCKET NUMBER: 32,944
REGISCOMMUNICATION INPORMATION:
TELECOMMUNICATION INPORMATION:
TELECOMMUNICATION INPORMATION:
TELECOMMUNICATION INPORMATION:
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APPLICANT: Villeponteau, Bryant
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Julli
APPLICANT: Peng, Julli
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STRET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                          FILING DATE: 07-JUN-1995
CLASSIFICATION: 435
PRIOR APPLICATION 10ATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
ATORNEY AGENT: 1NFORMATION:
NAME: STOTELLA, JOHN R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-000830US
TELEPAN: (415) 576-0200
TELEPAN: (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4.4%; Scc...
100.0%; Pred. No. c...
0; Mismatches
APPLICATION NUMBER: US/08/482,115B FILING DATE: 07-JUN-1995
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ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 7, Application US/08660678A Patent No. 5837857
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Best Local Similarity 100.
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; MOLECULE TYPE: DNA
US-08-482-115B-7
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US-08-660-678A-7/c
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Gaps
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Patent No. 5846723
GENERAL INFORMATION:
APPLICANT: Kim, Mam Woo
APPLICANT: Win, Fred
APPLICANT: Weinley, James T.
APPLICANT: Weinrich, Scott L.
TITLE OF INVENTION: Methods for Detecting the RNA Component of TITLE OF INVENTION: Telomerase
NUMBER OF SEQUENCE: 26
CORRESPONDENCE ADDRESS:
ADDRESSEE: TOWNSEND and TOWNSEND and CREW LLP
STREET: TWO Embarcadero Center, 8th Floor
CITY: San Francisco
STATE: California
COUNTRY: USA
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83;
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TITLE OF INVENTION: Telomerase
NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
ADDRESSEE: TOWNSEND and TOWNSEND and CREW LLP
STREET: TWO Embarcadero Center, 8th Floor
CITY: San Francisco
STATE: California
COMPUTR: USA
ZIP: 9411-3834
COMPUTR: HAPE: Floppy disk
COMPUTR: ISM PC compatible
COMPUTR: ISM PC compatible
COMPUTR: Floppy disk
COMPUTR: ISM PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/770,565
FILING DATE: 20-DEC-1996
CLASSIFICATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 32,944
REGISTRATION NUMBER: 32,94
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APPLICATION NUMBER: US/08/770,565 FILING DATE: 20-DEC-1996 CLASSIFICATION: 435
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     361 AGGCCGCAGGAAGAGGAACG 380
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CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity 100.
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-08-770-565-6/c
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Patent No. 5846723
GENERAL INFORMATION:
APPLICANT: Kim, Nam Woo
APPLICANT: Kealey, James T.
APPLICANT: Realey, James T.
APPLICANT: Weinrich, Scott L.
TITLE OF INVENTION: Methods for Detecting the RNA Component of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
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Pred. No. 83;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: CALLLULLIAGE
CONFUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/660,678A
FILING DATE: 05-JUN-1996
CLASSITCATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/272,102
FILING DATE: 07-JUL-1994
ATTORNEY/AGENT INFORMATION:
NAME: Stochalla, JOHN R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-000811US
TELECOMMUNICATION NUMBER: 015389-000811US
TELECOMMUNICATION NUMBER: 015389-000811US
                                                                                                                                                                                                                                                            Sequence 29, Application US/08660678A

Patent No. 5837857

GENERAL INFORMATION:

APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junli

APPLICANT: Andrews, William H.

TITLE OF INVENTION: Mammalian Telomerase

NUMBER OF SEQUENCES: 30

CORRESPONDENCE ADDRESS:

ADDRESSEE: Townsend and Townsend and Crew LLP

STREET: Two Embarcadero Center, Eighth Floor

CITY: San Francisco

SCALIFORMATION:

CONTRESS OF SECUENCES:

CITY: SAN Francisco

SCALIFORMATION:

CONTRESS OF SECUENCES:

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100.0%; Pred. No. co...
0; Mismatches
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                             20 TIGITANCCTAACTGAG 1
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TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 29:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 100.
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; TOPOLOGY: linear; MOLECULE TYPE: RNA US-08-660-678A-29
                                                                                                                                                                                                 RESULT 125
US-08-660-678A-29/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-08-770-565-3/c
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Gaps
                                                                                                                                                      GENERAL INFORMATION:
APPLICANT: Kim, Nam Woo
APPLICANT: Kim, Nam Woo
APPLICANT: Kaley, James T.
APPLICANT: Realey, James T.
APPLICANT: Weinrich, Scott L.
TITLE OF INVENTION: Telomerase
NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
ADDRESSEE: TOWNSEND and TOWNSEND and CREW LLP
STREET: Two Embarcadero Center, 8th Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
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                                                                                                                                                                                                                                                                                                                                                                                              STATE: CALILLIANCE
COMPUTRY: USA
ZIP: 9411-3834

COMPUTRY: USA
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTRI: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN DATA:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/770,565
FILING DATE: 20-DEC-1996
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERRACE/DOCKET NUMBER: 015389-002300US
TELEPAX: 415-576-0200
TELEPAX: 415-576-0300
TELEFAX: 415-576-0300
TELEFAX: 415-576-0300
TELEFAX: 20 Dase pairs
290 CTGCCACCGCGAAGAGTTGG 309
                                                                                                                        ; Sequence 13, Application US/08770565; Patent No. 5846723
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      159 TCTAGAGCAAACAAAAATG 178
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 6, Application US/08485778
Patent No. 5876979
GENERAL INFORMATION:
                        20 CTGCCACCGCGAAGAGTTGG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20 TCTAGAGCAAACAAAATG 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
Matches 20; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; TOPOLOGY: linear; MOLECULE TYPE: DNA US-08-770-565-13
                                                                                     RESULT 129
US-08-770-565-13/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
PILLING DATE: US/08/770,565
FILING DATE: 20-DEC-1996
         32,944
PRR: 015389-002300US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION: 435
ATTORNEY/ASTRI INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REEREROE/DOCKET NUMBER: 015389-002300US
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                          GTCAG 319
         REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 01530
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-576-0200
TELEPAX: 415-576-0300
INFORMATION FOR SEQ ID NO: 6: SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                          20 GAAGAGTIGGGCTCTGTCAG 1
                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 100.
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: 415-576-0300
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity 100.
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                        300 GAAGAGTIGGGCTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: nucleic acid
STRANDEDNESS: sing
                                                                                                                                                                                             ; TOPOLOGY: linear; MOLECULE TYPE: DNA US-08-770-565-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-08-770-565-7
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APPLICANT: Avilion, Ariel Athena
APPLICANT: Feng, Junil
APPLICANT: Feng, Junil
APPLICANT: Greider, Carol
APPLICANT: Marhuenda, Maria Antonia Blasco
APPLICANT: Villeponteau, Bryant
TITLE OF INVENTION: RNA COMPONENT OF TELOMERASE
NUMBER OF SEQUENCES: 45
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/485,778
FILING DATE: 07-JE-1995
CLASSFEICATION: 435
RICH APPLICATION DATA:
APPLICATION NUMBER: US 08/387,524
FILING DATE: 13-FEB-1995
PRIOR APPLICATION NUMBER: US 08/330,123
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
ATYORNEY/AGENT INPORMATION:
NAME: GRANAHAMIER: 32,227
REGISTRATION NUMBER: 32,227
REFERENCE/POCKET NUMBER: 32,227
REFERENCE/POCKET NUMBER: CSHL94-05A4
TELEPHONE: 617-661-6240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER FRADABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/472,802C
FILING DATE: 07-JUN-1995
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
PRIOR APPLICATION DATA:
PRILOR APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/30,123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               | Second | S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2 GGTTGCGGAGGGTGGCCTG 21
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ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: 9
SEQUENCE CHARACTERISTICS:
TRUGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; TOPOLOGY: linear
US-08-485-778-9
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US-08-472-802C-8/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Andrews, William H.
APPLICANT: Andrews, William H.
APPLICANT: Feng, Juni;
APPLICANT: Feng, Juni;
APPLICANT: Funk, Walter
APPLICANT: Funk, Walter
APPLICANT: Walter
APPLICANT: Warhuenda, Maria Antonia Blasco
APPLICANT: Villeponteau, Bryant
ITITE OF INVENTION: RNA COMPONENT OF TELOMERASE
NUMBER OF SEQUENCES: 45
CORRESPENDENCE ADDRESS:
ADDRESSE: Hamilton, Brook, Smith & Reynolds, P.C.
STREET: Two Militia Drive
CITY: Lexington
STATE: MA
COUNTRY: US
ZIP.
            CORRESPONDENCE ADDRESS:
ADDRESSE: Hamilton, Brook, Smith & Reynolds, P.C.
STREET: Two Militia Drive
CITY: Lexington
STRYE: MA
STRYE: MA
STRYE: US
ZIP: 02173
                                                                                                                                                                                                                                                                                   COMPUTER EADABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/485,778
FILING DATE: 07-JE-1995
CLASSIFICATION: 435
PRIOR APPLICATION ADATA:
APPLICATION NUMBER: US 08/387,524
FILING DATE: 13-FEB-1995
PRIOR APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-CT-1994
FILING DATE: 27-CT-1994
FILING DATE: 07-JUL-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TTTGTCTAACCCTAACTGAG 60
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: 617-861-9540
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 100.
Matches 20; Conservative
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US-08-485-778-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-08-485-778-9/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STALL
COMPURY: USA
ZIP: 9411-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/472,802C
FILING BATE: 07-JUN-1995
FILING BATE: 07-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 133
US-08-472-802C-41/C
; Sequence 41, Application US/08472802C
; Patent No. 5958680
; GENERAL INFORMATION:
    APPLICANT: Villeponteau, Bryant
    APPLICANT: Andrews, William H.
    TITLE OF INVENTION: Mammalian Telomerase
; NUMBER OF SEQUENCES; 44
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Townsend and Townsend and Crew
        TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEPHONE: (415) 576-0200
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDENNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 07-UN-1995
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-UL-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M.
REGISTRATION NUMBER: 30,223
REFENCE/DOCKET NUMBER: 33,223
REFENCE/DOCKET NUMBER: 15389-00082
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                     2 GGTTGCGGAGGGTGGGCCTG 21
REGISTRATION NUMBER: 30,223
                                                                                                                                                                                                                TOPOLOGY: linear MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                      Best Local Similarity 100.
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: Two concerts CITY: San Francisco cTATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; MOLECULE TYPE: RNA
US-08-472-802C-41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        linear
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4.4%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 83;

Query Match Best Local Similarity

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Gaps
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Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/833,377
FILING DATE: 04-APR-1997
CLASSIFICATION: 435
PRIOR APPLICATION NUMBER: US/08/510,736
FILING DATE: 04-APR-1995
ATFORNEY/AGENT INPORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 37,040
TELEPRA: (415) 576-0200
TELEPRA: (415) 576-0200
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LEMOTH: 20 base pairs
LUMBER: LEMOTH: 20 base pairs
LUMBER: LEMOTH: 20 base pairs
LUMBER: LEMOTH: 20 base pairs
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Townsend and Townsend and Crew LLP
                                                                                                                                                                                                                                                                                                                                                                                                                        JUMNSEND AUWISEND AND TOWNSEND AND Crew LL CITY: San Francisco STATE: California COUNTRY: USA
                                                                                                                                                                                        Sequence 7, Application US/08833377

Patent No. 5968506

GENERAL INFORMATION:

APPLICANT: Meinrich, Scott L.

APPLICANT: Atkinson III, Edward M.

APPLICANT: Lichtsteiner, Serge P.

APPLICANT: Pruzan, Ronald A.

APPLICANT: Pruzan, Ronald A.

APPLICANT: Realey, James T.

TITLE OF INVENTION: Purified Telomerase

NUMBER OF SEQUENCES: 15

CORRESPONDENCE ADDRESS:
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: -
LOCATION: 1..20
OTHER INFORMATION: /note= "Oligo 14ab"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 135
US-08-520-550A-6/c
; Sequence 6, Application US/08520550A
; Patent No. 6013468
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        361 AGGCCGCAGGAAGAGGAACG 380
                                            41 TTTGTCTAACCCTAACTGAG 60
                                                                                   20 rrrercraacceraacreag 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20 AGGCCGCAGGAAGAGGAACG
20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity 100.
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRANDEDNESS:
                                                                                                                                                                            US-08-833-377-7/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-08-833-377-7
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                                                                                                                                                       RESULT 134
Matches
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Sequence 13, Application US/08630019A
Fatent No. 6015710
GENERAL INFORMATION:
APPLICANT: Wright, Woodring E.
APPLICANT: Platyszek, Mieczyslaw A.
APPLICANT: Platyszek, Mieczyslaw A.
APPLICANT: Orerey, David
APPLICANT: Correy, David
APPLICANT: Correy, David
APPLICANT: Correy, David
APPLICANT: Posting E.
APPLICANT: Posting E.
APPLICANT: Posting E.
APPLICANT: Profession of Mammalian Telomerase by
TITLE OF INVENTION: Modulation of Mammalian Telomerase by
TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 46
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 1; Length 20;
83;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Indela
                                                                                                            COUNTRY: US
ZIP: 02173
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/520,550A
FILING DATE: 29-AUG-1995
CLASSIFICATION NUMBER: US 08/387,524
FILING DATE: 13-FEB-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/337,524
FILING DATE: 17-CCT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
PRIOR APPLICATION NUMBER: US 08/272,102
FILING DATE: 37-CCT-1994
PRIOR APPLICATION NUMBER: US 08/272,102
FILING DATE: 17-CT-1994
TILING DATE: 17-CT-1994
TELECOMMUNICATION NUMBER: CSHL94-05A3B
TELECOMMUNICATION INPERMATION:
TELEBRONE: 617-861-5940
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE: HARACTERISTICS:
LENGTH: 20 base pairs
3: Hamilton, Brook, Smith & Reynolds, P.C.
Two Militia Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 4.4%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 83; Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 94111-3834

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2 GGTTGCGGAGGGTGGGCCTG 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: single
                STREET: Two ...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; TOPOLOGY: linear
US-08-520-550A-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-08-630-019A-13/c
     ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ð
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             4.4%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 83; 0; Indels tive 0; Mismatches 0; Indels
                                                                      APPLICANT: Funk, Walter
APPLICANT: Greider, Carol
APPLICANT: Marhuenda, Maria A. B.
APPLICANT: Marhuenda, Maria A. B.
TITLE OF INVENTION: RNA Component of Telomerase
NUMBER OF SEQUENCES: 47
CORRESPONDENCE ADDRESS:
ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C.
STREET: Two Militia Drive
CITY: Lexington
                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: US
ZIP: 02173
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US 08/520,550A
FILING DATE: 29-AUG-1995
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/387,524
FILING DATE: 13-FEB-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
ATPORNEY/AGENT INPORMATION:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
ATTORNEY/AGENT INPORMATION:
APPLICATION NUMBER: 32,227
REGISTRATION NUMBER: 32,227
REGISTRATION NUMBER: 32,227
TELEFAX: 617-861-6240
INFORMATION FOR SEG ID NO: 6:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
INPORMATION FOR SEG ID NO: 6:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
ITYPE: nucleic acid
TYPE: nucleic acid
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Andrews, William H.
APPLICANT: Andrews, William H.
APPLICANT: Feng, Junii
APPLICANT: Feng, Junii
APPLICANT: Greider, Carol
APPLICANT: Marhuenda, Maria A. B.
APPLICANT: Willeponteau, Bryant
ATILE OF INVINION: RNA Component of Telomerase
NUMBER OF SEQUENCES: 47
CORRESPONDENCE ADDRESS:
Andrews, William H.
Avilion, Ariel A.
Feng, Junli
Funk, Walter
Greider, Carol
Marhuenda, Maria A. B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 9, Application US/08520550A
Patent No. 6013468
GENERAL INFORMATION:
APPLICANT: Andrews, William H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          41 TITGICTAACCCTAACTGAG 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity 100.
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                            CITY: Lexir
STATE: MA
COUNTRY: U
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-08-520-550A-9/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-08-520-550A-6
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Gape

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iENGTH: 20 base pairs
TYPE: mucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
NOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "peptide nucleic acid (PNA),
DESCRIPTION: where (deoxy(ribose-phosphate linkages are replaced by DESCRIPTION: N-(2-aminoethy1)glycine units linked to nucleotide bases via DESCRIPTION: glycine amino N through a methylenecarbonyl linker"
US-08-838-545-41
                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                            ..
0
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                                                                                                                                                                                                                                                                                                 Query Match
4.4%; Score 20; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 83;
Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junli
APPLICANT: Funk, Walter
APPLICANT: Funk, Walter
APPLICANT: Andrews, William H.
ITILE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/998,443 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                015389-000811US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4.4%; Score 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRICA APPLICATION DATA:
APPLICATION NUMBER: US/08/660,678
FILING DATE: 05-UN-1996
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-0CT-1994
PRIOR APPLICATION NUMBER: US 08/272,102
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-UL-1994
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-0008
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
COMPUTER: FLOUDY disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                               46 CTAACCCTAACTGAGAAGGG 65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-08-998-443-7/c
; Sequence 7, Application US/08998443
; Patent No. 6054575
                                                                                                                                                                                                                                                                                                                                                                                                                                                               20 CTAACCCTAACTGAGAAGGG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-08-998-443-7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 139
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               /desc = "peptide nucleic acid (PNA), where (deoxy)ribose-phosphate linkages are replaced by N-(2-aminoethyl)glycine units linked to nucleotide bases via glycine amino nitrogen through a methylenecarbonyl linker"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Shay, Jerry W.
APPLICANT: Shay, Jerry W.
APPLICANT: Wright, Woodring E.
APPLICANT: Piatyszek, Mieczyslaw A.
APPLICANT: Orecy, David R.
APPLICANT: No. 6046307ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by
TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SURGENESS: 60
CORRESPONDENCE ADDRESS:
ADDRESSE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SIATE: CALIFOLITIA

SIATE: CALIFOLITIA

ZIP: 94111-3834

COMPUTER READABLE FORM:
MEDLUW TYPE: FOLDPY disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN RElease #1.0, Version #1.30

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/838,545
FILING DATE: 09-APR-1997
FILING DATE: 09-APR-1997
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INFORMATION:
NAME: SLOCATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 31,944
REFERENCE/DOCKET NUMBER: 015389-001610US
TELEPHONE: (415) 576-0200
TELEFPHONE: (415) 576-0200
INFORMATION FOR SEQ ID NO: 41:
             ATTORNAY ON NUMBER: US/08/630,019A
PILING DATE: 09-UNN-1996
CLASSIFICATION: 536
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-001600US
TELECOMMUNICATION INFORMATION:
TELEFRANCE/TOCKET NUMBER: 015389-001600US
TELEFRANCE/TOCKET NUMBER: 015389-001600US
TELEFRANCE/TOCKET NUMBER: 015389-001600US
TELEFAX: (415) 576-0300
TELEFAX: (415) 576-0300
TELEFAX: (415) 576-0300
TELEFAX: CARRACTERISTICS: 13:
SEQUENCE CHARACTERISTICS: LENGTH: 20 base pairs
TYPEN: nucleic acid
STRANDENESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
4.4%; Score 20; DB 1;
Best Local Similarity 100.0%; Pred. No. 83;
Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                             other nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 41, Application US/08838545 Patent No. 6046307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              46 CTAACCCTAACTGAGAAGGG 65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20 craacccraacreagaage 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20; Conservative
     CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                    linear
                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: 1in
MOLECULE TYPE:
DESCRIPTION:
DESCRIPTION:
DESCRIPTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 138
US-08-838-545-41/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-630-019A-13
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Gaps
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83;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/998,443 FILING DATE:
                                                   APPLICANT: Funk, Walter
APPLICANT: Funk, Walter
APPLICANT: Andrews, William H.
ITILE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLE
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/660,678
FILING DATE: 05-UN-1996
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-CCT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-UL-1994
ATTORNEY/AGENT INFORMATION:
NAME: SECREIA, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-000811US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 29:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 ABBE pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 7, Application US/09060523;
Patent No. 6258535;
GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junli
APPLICANT: Feng, Junli
APPLICANT: Fank, Walter
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 25;
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 4.4%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 83; Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                           PC-DOS/MS-DOS
         Villeponteau, Bryant
Feng, Junli
Funk, Walter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2 GGTTGCGGAGGGTGGGCCTG 21
                                                                                                                                                                                                                                                                COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IRN PC COMPALIALE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20 GerrecesAccerces 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                         OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; TOPOLOGY: linear; MOLECULE TYPE: RNA US-08-998-443-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-09-060-523-7/c
         APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ò
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                                    Gaps
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4.4%; Score 20; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 83;
Matches 20; Conservative 0; Mismatches 0; Indels
                                    0; Indels
                                                                                                                                                                                             RESULT 140

US-08-999-443-26/C

i Sequence 26, Application US/08998443

i Patent No. 6054575

i GENERAL INFORMATION:

APPLICANT: Feng, Junil

APPLICANT: Funk, Walter

APPLICANT: Funk, Walter

APPLICANT: Andrews, William H.

TITLE OF INVENTION: Mammalian Telomerase

NUMBER OF SEQUENCES: 30

CORRESPONDENCE ADDRESS:

ADDRESSEE: Townsend and Townsend and Crew LLP

STREET: Two Embarcadero Center, Eighth Floor

CITY: San Francisco

STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: USA
ZIP: 94111-3834
COMPUTER REDABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATCHITIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/998,443
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIORESTECATION NOMBER: US/US/959,443
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/660,678
FILING DATE: 05-JUN-1996
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
ATORNEY AGENT INFORMATION:
NAME: SCOFELLA, JOHN R.
REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 26:
SEQUENCE CHARACTERISTICS:
LENGTH: 120 ABBE pairs
100.0%; Pred. ...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          41 TTTGTCTAACCCTAACTGAG 60
                                                                                 2 GGTTGCGGAGGGTGGGCCTG 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20 TTTGTCTAACCCTAACTGAG 1
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         Best Local Similarity 100. Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-08-998-443-26
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ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor STRY: San Francisco STATE: California

ZIP: 94111-3834

COUNTRY:

US-08-998-443-29/c ; Sequence 29, Application US/08998443 ; Patent No. 6054575 ; GENERAL INFORMATION:

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other nucleic acid (PNA), desc = "peptide nucleic acid (PNA), where (deoxy/ribose-phosphate linkages are replaced by N-(2-aminoethy)2ycine units linked to nucleotide bases via glycine amino N through a methylenecarbonyl linker"
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MEDIUM TYBE FLOPDY disk
COMPUTER READABLE FORM:

MEDIUM TYBE: FLOPDY disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: DEACHLIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/580,517
FILING DATE: 25-MAY-2000
CLASSIFICATION DATA:
APPLICATION NUMBER: 08/330,123
FILING DATE: cunknown>
ATTORNEY/AGENT INFORMATION:
NAME: SMITCH, William M
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 15389-000810
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES: 25
CORRESPONDENCE ADDRESS:
ADDRESSER: Townsend and Townsend Khourie and Crew
STREET: 379 Lytton Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
4.4%; Score 20; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 83;
Matches 20; Conservative 0; Mismatches 0; Indels
          APPLICATION NUMBER: US 08/838,545
FILING DATE: 09-APR-1997
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-ARR-1996
ATTORNEY/AGENT INFORMATION:
NAME: SFORTIA, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-001610US
TELEPHONE: (415) 576-0200
TELEPHONE: (415) 576-0200
TELEPEAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 41:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDENNES: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-09-580-517-4/c
; Sequence 4, Application US/09580517
; Patent No. 6320039
; GENERAL INFORMATION:
APPLICANT: VILLEPONTEAU, Bryant FENG, Juni;
FENG, Juni;
ANDREWS, William H.
TITLE OF INVENTION: HUMAN TELOMERASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: (415) 326-2400
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        46 CTAACCCTAACTGAGAAGGG 65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20 craacccraacreagaage 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Palo Alto
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                   linear
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MOLECULE TYPE:
DESCRIPTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DESCRIPTION:
DESCRIPTION:
DESCRIPTION:
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US-09-349-532-41/C

Sequence 41, Application US/09349532

Patent No. 6224650

GENERAL INFORMATION:
APPLICANT: Shay, Jerry W.
APPLICANT: Piatyszek, Mieczyslaw A.
APPLICANT: Orevy, David R.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STREET: Two Embarcadero Center, Eighth Floor
STREET: Tan Embarcadero Center, Eighth Floor
STREET: Talifornia
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COMPUTER READABLE FORM:

MEDIUM TYRE: Floppy disk
COMPUTER: IBM FC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US 08/60,523
FILING DATE: 14-APR-1998
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/60,678
FILING DATE: US 08/330,123
FILING DATE: 20-CCT-1994
FRICH APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
ATTORNEY/AGENT INFORMATION:
NAME: STORELLS, JOHN R.
REFERENCE/DOCKET NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 32,946
TELEPHONE: (415) 576-0300
TELEPHONE: (415) 576-0300
TELEPHONE: GLOSON DATES
TELEPHONE (415) 576-0300
TELEPHONE: CLOSON DATES
TELEPHONE (415) 576-0300
TELEPHONE: CLOSON DATES
TELEPHONE (415) 576-0300
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
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COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

SOFTWARE: PATENTIN Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

FILING DATE:

CLASSIFICATION:

PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2 GGTTGCGGAGGGTGGGCCTG 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20 GGTTGCGGAGGTGGGCCTG 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: DNA
US-09-060-523-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ద
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4.4%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 83; tive 0; Mismatches 0; Indels
                                                                                                                                       Length 20;
                                                                                                                                                                            0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: US
ZIP: 94301
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/330,123
FILING DATE: «Unknown:
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 15389-000810
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION: (415) 326-2400
                                                                                                                                     Query Match
4.4%; Score 20; DB 1;
Best Local Similarity 100.0%; Pred. No. 83;
Matches 20; Conservative 0; Mismatches (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/580,517
FILING DATE: 25-May-2000
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ANDREWS, William H.
TITLE OF INVENTION: HUMAN TELOMERASE
NUMBER OF SEQUENCES: 25
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: DNA
SEQUENCE DESCRIPTION: SEQ ID NO: 7:
                                         ) MOLECULE TYPE: RNA
SEQUENCE DESCRIPTION: SEQ ID NO: 4:
US-09-580-517-4
                                                                                                                                                                                                                                                                                                                                  US-09-580-517-7/c

| Sequence 7, Application US/09580517
| Patent No. 6320039
| GENERAL INFORMATION:
| APPLICANT: VILLEPONTEAU, Bryant
                                                                                                                                                                                                                    41 TTTGTCTAACCCTAACTGAG 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2 GGTTGCGGAGGGTGGGCCTG 21
                                                                                                                                                                                                                                                 20 TTTGTCTAACCCTAACTGAG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FELEFAX: (415) 326-2422
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20 GGTTGCGGAGGGTGGGCCTG 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: nucleic acid
STRANDEDNESS: single
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                        FENG, Junli
FUNK, Walter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO: 7
                                       TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity
Matches 20; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-09-580-517-7
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                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: BY PC compatible

COMPUTER: BY PC compatible

COMPUTER: BY PC compatible

SOFTWARE: PatentIn Release #1.0, Version #1.30

CURRENT APPLICATION NUMBER: US/09/057,351

FILING DATE: 08-APR-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Townsend and Townsend and Crew LLP
                                                                                                                                                                                            STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco STATE: California COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE:

PRIOR APPLICATION: 435

PRIOR APPLICATION DATA:

APPLICATION UNDER:

FILING DATE: 07-UL-1994

PRIOR APPLICATION DATA:

APPLICATION WUMBER: US 08/330,123

FILING DATE: 27-OCT-1994

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/472,802

FILING DATE: 07-UN-1995

ATTORNEY AGENT INFORMATION:

NAME: Storella, John R.

REGISTATION NUMBER: 32,944

REFERENCE/DOCKET NUMBER: 015389-000821US

TELECOMMUNICATION INFORMATION:

TELEFAM: (415) 576-0200

TELEFAM: (415) 576-0300

INFORMATION FOR SEQ ID NO: 75 SEQUENCE CHARACTER/STICS:
US-09-057-351-7/c

Sequence 7, Application US/09057351

Patent No. 6548049

APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junil
APPLICANT: Funk, Walter
APPLICANT: Funk, Walter
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 42

CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Crew
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junil
APPLICANT: Funk, Walter
APPLICANT: Funk, Walter
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 100.0%; Fred. No. 83;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-09-057-351-40/c
; Sequence 40, Application US/09057351
; Patent No. 6548298
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2 GGTTGCGGAGGGTGGGCCTG 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; MOLECULE TYPE: DNA US-09-057-351-7
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RESULT 146

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; Sequence 60, Application US/08838545;
; Patent No. 6046307;
; RENERAL INFORMATION:
    APPLICANT: Wight, Woodring E. APPLICANT: Piatyszek, Mieczyslaw A. APPLICANT: Orey, David R. APPLICANT: No. 6046307ton, James C. TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Peptide Nucleic Acids NUMBER OF SEQUENCES: 60 CORRESPONDENCE ADDRESS: ADDRESSE: Townsend and Crew LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1; Length 19; 93;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SCFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/088.545
FILING DATE: 09-APR-1997
CLASSIFICATION 536
PRIOR APPLICATION NUMBER: US 08/630,019
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTONNEY/AGENT INPORMATION:
NAME: SCORELLA, JOHN R.
REGISTRATION NUMBER: 32,944
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco STATE: California COUNTR: USA ZIP: 94111-3834 COMPUTER READABLE FORM: MEDLUM TYPE: Floppy disk COMPUTER: IBM PC COMPALIBLE OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                    ATTORNEY AGENT INFORMATION:

NAME: Storella, John R.
REGISTATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-002300US
TELEFONE. 1415-576-0200
TELEFAX: 415-576-0300
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4.2%; Scor.
100.0%; Pred. No. 20.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REFERENCE JOCKET NUMBER: 04,944
REFERENCE JOCKET NUMBER: 015389-001610US
TELECOMMUNICATION INFORMATION:
TELEPRONE: (415) 576-0200
INFORMATION FOR SEQ ID NO: 60:
SEQUENCE CHARACTERISTICS:
LENGTH: 19 base pairs
TENGTH: 19 base pairs
TENGTH: 19 base pairs
TENGTH: 19 base pairs
148 CCACCGTTCATTCTAGAGC 166
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Best Local Similarity 100.0
Matches 19, Conservative
               CURRENT APPLICATION DATA:
                                     APPLICATION NUMBER:
FILING DATE: 20-DEC
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear; MOLECULE TYPE: DNA US-08-770-565-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRANDEDNESS:
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; Sequence 9, Application US/08770565
; Patent No. 5846723
; GENERAL INFORMATION:
APPLICANT: Wi, Krad
APPLICANT: Win Nam Woo
; APPLICANT: Fred
APPLICANT: Pruzan, Ronald
APPLICANT: Pruzan, Ronald
TITLE OF INVENTION: Methods for Detecting the RNA Component of
TITLE OF INVENTION: Telomerase
; TITLE OF INVENTION: Telomerase
; NUMBER OF SEQUENCES: 26
; CORRESPONDENCE ADDRESS: 26
; ADDRESSEE: TOWNSEND and TOWNSEND and CREW LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4.4%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 83; tive 0; Mismatches 0; Indels
                                                                                   COUNTRY: USA

ZIP: 9411-3934

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IB PC compatible
OPERATIOR SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US 08/09/057,351
FILING DATE: 08-APR-1994
CLASSIFICATION DATA:
APPLICATICATION DATA:
APPLICATICATION DATA:
APPLICATICATION DATA:
APPLICATICATION DATA:
APPLICATICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 07-JUL-1994
PRIOR APPLICATION NUMBER: US 08/472,802
FILING DATE: 07-JUN-1995
ATTORNEY/AGENT INFORMATION:
APPLICATION NUMBER: US 08/472,802
FILING DATE: 07-JUN-1995
ATTORNEY/AGENT INFORMATION:
AREFERENCE/DOCKET NUMBER: 015389-000821US
TELECOMMUNICATION NUMBER: 015389-000821US
TELEPHONE: (415) 576-0200
TELEPAX: (415) 576-0200
TELEPAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 40:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TTELEPAX: sinqle
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/WS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
            Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Two Embarcadero Center, 8th Floor
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Best Local Similarity 100.(
Matches 20; Conservative
                                San Francisco
California
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CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear MOLECULE TYPE: RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-770-565-9/c
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Gaps

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Best Local Similarity
Matches 19; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-09-018-125-2
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                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Shay, Jerry W.
APPLICANT: Wight, Woodring E.
APPLICANT: Pietyezek, Mieczyslaw A.
APPLICANT: Pietyezek, Mieczyslaw A.
APPLICANT: Corey, David R.
APPLICANT: No. 6294650ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerage by TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 60
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Town Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                              Query Match 4.2%; Score 19; DB 1; Length 19; Best Local Similarity 78.9%; Pred. No. 93; Matches 15; Conservative 4; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4.2%; Score 19; DB 1; Length 19; 78.9%; Pred. No. 93;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/NS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/349,532
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRICE APPLICATION DATA:
APPLICATION NUMBER: US 08/838,545
FILING DATE: 09-APR-1997
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INFORMATION:
NAMB: Storella, John R.
REGISTRATION NUMBER: 32,944
RECISTRATION NUMBER: 32,944
RELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 60:
SEQUENCE CHARACTERISTICS:
LENGTH: 19 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4; Mismatches
                                                                                                                                                                                                                                                                                                           Sequence 60, Application US/09349532
Patent No. 6294650
GENERAL INFORMATION
APPLICANT: Wright, Woodring E.
                                                                                                                                                                                          44 GTCTAACCCTAACTGAGAA 62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             44 GTCTAACCCTAACTGAGAA 62
; TOPOLOGY: linear
; MOLECULE TYPE: RNA (genomic)
US-08-838-545-60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: RNA (genomic)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: nucleic acid
STRANDEDNESS: sing
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Best Local Similarity
Matches 15, Conserva
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CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Sar
STATE: Ca
COUNTRY:
                                                                                                                                                                                                                                                                                              US-09-349-532-60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-09-349-532-60
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RESULT 151

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                                                                        APPLICANT: Silverman, Robert H.
APPLICANT: Silverman, Robert H.
APPLICANT: Silverman, Robert H.
APPLICANT: Kondo, Seiji
APPLICANT: Cowell, John K.
APPLICANT: Cowell, John K.
APPLICANT: Torrence, Paul F.
TITLE OF INVENTION: RNASE L ACTIVATORS AND ANTISENSE OLIGONUCLEOTIDES
TITLE OF INVENTION: EFFECTIVE TO TREAT TELOMERASE-EXPRESSING MALIGNANCIES
FILE REPERENCE: 8656-022
CURRENT APPLICATION NUMBER: US/09/018,125A
CURRENT FILING DATE: 1999-02-03
EARLIER APPLICATION NUMBER: 60/044,507
EARLIER FILING DATE: 1997-04-21
NUMBER OF SEQ ID NOS: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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100.0%; Pred. No. 93;
iive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: Description of Artificial Sequence: OTHER INFORMATION: oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER PATABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC Compatible
COPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/833,377
FILING DATE: 04-APR-1997
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco STATE: Califernia COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: 015389-001110US TELECOMMUNICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 14, Application US/08833377
; Sequence 14, Application US/08833377
; Patent No. 5968506
; GENERAL INFORMATION:
    APPLICANT: Weinrich, Scott L.
    APPLICANT: Atkinson III, Edward M.
    APPLICANT: Lichtsteiner, Serge P.
    APPLICANT: Vasserot, Alain P.
    APPLICANT: Pruzan, Ronald A.
    APPLICANT: Kealey, James T.
    TITLE OF INVENTION: Purified Telomerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 04-APR-1997
CLASSIFICATION: 435
FICK PAPLICATION DATA:
APPLICATION NUMBER: US 08/510,736
Sequence 2, Application US/09018125A Patent No. 6468983 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            76 GIGCITITGCICCCCCCCC 94
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 04-AUG-1995
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: Patentin Ver. 2.0 SEQ ID NO 2 LENGTH: 19 TYPE: DNA ORGANISM: Artificial Sequence
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San Francisco
California
    linear
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Best Local Similarity
                  MOLECULE TYPE:
DESCRIPTION:
DESCRIPTION:
DESCRIPTION:
DESCRIPTION:
DESCRIPTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DESCRIPTION:
DESCRIPTION:
    TOPOLOGY:
                                                                                                                         US-08-838-545-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-09-349-532-9
                                                                                                                                                                 Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Wright, Woodring E.
APPLICANT: Wright, Woodring E.
APPLICANT: Wright, Woodring E.
APPLICANT: Piatyszek, Mieczyslaw A.
APPLICANT: Corey, David R.
APPLICANT: No. 6046307ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 60
CORRESPONDENCE ADDRESS:
ADDRESSEE: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                           /mod base= OTHER
/note= "N = 5' biotinylated cytidine'
                                                                                                                                                                                                                                                                                                                                                                                                         Length 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Indels
                                                                                                                                                                                                                                                                                 FEATURE:

NAME/KEY:

LOCATION: 1..20

GTHER INFORMATION: /note= "biotinylated Oligo 14ab"
US-08-833-377-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OPERATION SYSTEM: PC-DUS, PS-DUS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/838,545
FILING DATE: 09-APR-1997
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INFORMATION:
NAME: SLOCATION NUMBER: 015389-001610US
FELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                       Score 19; DB 1;
Pred. No. 99;
                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; Pred. ...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-08-838-545-9/c; Sequence 9, Application US/08838545; Patent No. 6046307; Batent INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        361 AGGCCGCAGGAAGAGGAAC 379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20 AGGCCGCAGGAAGAGAAC 2
  (415) 576-0200
              TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 14
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                     NAME/KEY: modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 18 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
                                                                                                                                                                                                                        LOCATION: 1
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                        TOPOLOGY: linear
MOLECULE TYPE: DNA
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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other nucleic acid (PNA), desc = "peptide nucleic acid (PNA), where (deoxy/ribose-phosphate linkages are replaced by N.(2-aminoethyl)dlycine units linked to nucleotide bases via glycine amino N through a methylenecarbonyl linker"
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/desc = "peptide nucleic acid (PNA),
where (deoxy/ribose-phosphate linkages are replaced by
N.(2-aminoechy) alycine units linked to nucleotide bases via
glycine amino N through a methylenecarbonyl linker"
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                                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: Shay, Jerry W.
APPLICANT: Wright, Woodring E.
APPLICANT: Partysek, Micczyslaw A.
APPLICANT: Partysek, Micczyslaw A.
APPLICANT: Corey, David R.
APPLICANT: Corey, David R.
APPLICANT: Corey, David R.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 60
CORRESPONDENCE ADDRESS:
ADDRESSER: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                             DB 1; Length 18; 1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 18;
                                                                                                                                                                                                                                               0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: USA

COUNTRY: USA

COMPUTER: READABLE FORM:
MEDIUM TYPE: FIDEPY disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: PATENTING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/09/349,532
FILING DATE:
                                                                                                                                                                                             4.0%; Score 18; DB 100.0%; Pred. No. 1e-tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4.0%; Score 18; 100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION:
PRIOR APPLICATION DATA:
PRIOR APPLICATION UNDBER:
PILING DATE: 09-APR-1997
PILING DATE: 09-APR-1966
ATTOCNEV/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
FREISTRATION NUMBER: 32,944
                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-09-349-532-9/c
; Sequence 9, Application US/09349532
; Patent No. 6294650
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REFERENCE/DOCKET WUMBER: 015.3 TELECOMMUNICATION: TELECHONE: (415) 576-0200 TELEFAX: (415) 576-0300 INPORMATION FOR SEQ ID NO: 9: SEQUENCE FARACTERISTICS: LENGTH: 18 base pairs
                                                                                                                                                                                                                                                                                                       48 AACCCTAACTGAGAAGGG 65
                                                                                                                                                                                                                                                                                                                                                      18 AACCCTAACTGAGAGGG 1
                                                                                                                                                                                                                      Best Local Similarity 100.
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      nucleic acid
EDNESS: single
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3.6%; Score 16.4; DB 1; Length 18; 94.4%; Pred. No. 1.4e+02; tive 0; Mismatches 1; Indels
Human Telomerase Catalytic Subunit 727
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OTHER INFORMATION: /note= "antisense hTRT molecule"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OPERATING SYSTEM: PC-DOS/MS-DOS
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/974,549A
FILING DATE: 19-NOV-1997
CLASSIFICATION NUMBER: US/08/974,549A
FILING DATE: 19-NOV-1997
CLASSIFICATION NUMBER: US/08/974,643
APPLICATION NUMBER: US/08/44,419
FILING DATE: 01-OCT-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/44,419
FILING DATE: 18-APR-1997
PRIOR APPLICATION NUMBER: US/08/46,017
FILING DATE: 25-APR-1997
PRIOR APPLICATION NUMBER: US/08/91,937
PRIOR APPLICATION NUMBER: US/08/91,937
PRIOR APPLICATION NUMBER: US/08/91,312
PRIOR APPLICATION NUMBER: US/08/91,312
PRIOR APPLICATION NUMBER: US/08/91,312
PRIOR APPLICATION NUMBER: US/08/91,312
FILING DATE: 14-AUG-1997
PRIOR APPLICATION NUMBER: US/08/912,951
FILING DATE: 14-AUG-1997
PRIOR APPLICATION NUMBER: US/08/915,503
FILING DATE: US/08/91
PRIOR APPLICATION NUMBER: US/08/91/17618
FILING DATE: US/08/91
PRIOR APPLICATION NUMBER: US/08/915,503
FILING DATE: US/08/91
PRIOR APPLICATION NUMBER: US/08/915,503
FILING DATE: US/08/91
ATPONDENT/AMPER: US/08/91
ATPONDENT/AMPERIANDENT/AMPERIANDENT/AMPERIANDENT/AMPERIANDENT/AMPERIANDENT/AMPERIANDENT/AMPERIANDENT/AMPERIANDENT/AMPERIANDENT/AMPERIANDENT/AMPERIANDENT/AMPERIANDENT/AMPERIANDENT/AMPERIANDENT/AMPERIANDENT/AMPERIANDENT/AMPERIANDENT/AMPERIANDENT/AMPERIANDENT/
TITLE OF INVENTION: Human Telomerase Catalytic & NUMBER OF SEQUENCES: 727
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Apple, Randolph Ted
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002610US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 543:
SEQUENCE CHARACTERISTICS:
LENGTH: 18 base pairs
                                                                                                                                                                                                                                                           COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Flopy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         149 CACCGTTCATTCTAGAGC 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 94.4'
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME/KEY: -
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           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Kim, Nam Woo
APPLICANT: Wu, Fred
APPLICANT: Wo, Fred
APPLICANT: Fred
APPLICANT: Fred
APPLICANT: Fred
APPLICANT: Mediaty, James T.
APPLICANT: Weinrich, Scott L.
TITLE OF INVENTION: Methods for Detecting the RNA Component of
TITLE OF INVENTION: Telomerase
NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
ADDRESSEE: TOWNSEND and TOWNSEND and CREW LLP
STREET: Two Embarcadero Center, 8th Floor
CITY: San Francisco
STATE: California
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           0; Indels
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CURRENT APPLICATION DATA:

APPLICATION NUMBER:

FILING DATE: 105/08/770,565

FILING DATE: 20-DEC-1996

CLASSIFICATION: 435

ATTONREY/AGENT INPORMATION:

NAME: Storella John R:

REGISTRATION NUMBER: 32,944

REGISTRATION INFORMATION:

TELERRENCE/DOCKET NUMBER: 32,944

REFERENCE/DOCKET NUMBER: 32,944

REFERENCE/OFFET NUMBER: 32,944

REFERENCE/OFFET NUMBER: 32,944

REFERENCE/OFFET NUMBER: 31,944

REFERENCE/OFFET NUMBER: 32,944

REFERENCE/OFFET NUMBER: 31,944

REFERENCE/OFFET NUMBER: 31,944

RESERRENCE/OFFET NUMBER: 31,944

RESERVER NUMBER: 31,94
       Mismatches
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2.08-97-4-549A-543/c
; Sequence 543, Application US/08974549A
; Patent No. 6166178
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE PORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                         Sequence 14, Application US/08770565
Patent No. 5846723
GENERAL INFORMATION:
APPLICANT: Kim, Nam Woo
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Lingner, Joachim
Nakamura, Toru
Chapman, Karen B.
Morin, Gregg B.
Harley, Calvin B.
Andrews, William H.
                                                                                  48 AACCCTAACTGAGAAGGG 65
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                                                                                                                         18 AACCCTAACTGAGAAGGG 1
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       18; Conservative
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STRANDEDNESS: single
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Best Local Similarity
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APPLICANT: Cech, Th
APPLICANT: Lingner,
APPLICANT: Nakamura
APPLICANT: Chapman,
APPLICANT: Morin, G
APPLICANT: Marley,
APPLICANT: Andrews,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              linear
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                                                                                                                                                                                                                                                                   RESULT 155
US-08-770-565-14/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-08-770-565-14
       Matches
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0; Gaps

US09436060A-16.rni.sl

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; IOCATION: 1..18 | IOCATION: 1..18 | SEQUENCE DESCRIPTION: SEQ ID NO: 543: US-09-402-181B-543
                                                                                                                                                                                                                                                                                                                         Nakamura, Toru
Chapman, Karen B.
Morin, Gregg B.
Harley, Calvin B.
Andrews, William H.
TITLE OF INVENTION: Human Telomerase Catalytic Subunit
NUMBER OF SEQUENCES: 633
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: INPE: Floppy disk

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/402,181B

FILING DATE: 29-5ep-1997

APPLICATION DATA:

APPLICATION DATA:

APPLICATION DATA:

APPLICATION NUMBER: US 08/844,419

FILING DATE: 10-0CT-1996

APPLICATION NUMBER: US 08/846,017

FILING DATE: 25-APR-1997

APPLICATION NUMBER: US 08/851,843

FILING DATE: 14-AUG-1997

APPLICATION NUMBER: US 08/911,312

FILING DATE: 14-AUG-1997

APPLICATION NUMBER: US 08/911,312

FILING DATE: 14-AUG-1997

APPLICATION NUMBER: US 08/912,951

FILING DATE: 14-AUG-1997

APPLICATION NUMBER: US 08/912,951

FILING DATE: 14-AUG-1997

APPLICATION NUMBER: WS 08/912,503

FILING DATE: 14-AUG-1997

APPLICATION NUMBER: WS 08/912,951

FILING DATE: 14-AUG-1997

APPLICATION NUMBER: WS 08/912,951

FILING DATE: 14-AUG-1997

APPLICATION NUMBER: WS 08/912,951
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INPORMATION:
NAME: Ausenbus, Scott L.
REGISTRATION NUMBER: 42,271
REFERENCE/DOCKET NUMBER: 015389-002620US
TELECOMMUNICATION INFORMATION:
                                                                                             RESULT 157
US-09-402-181B-543/C
; Sequence 543, Application US/09402181B
; Patent No. 6610839
; GENERAL INFORMATION:
; APPLICANT: Cech, Thomas R.
Lingner, Joachim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
18 CACCCTTCATTCTAGAGC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 18 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO: 543;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: San Francisco
STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: DNA
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Gaps
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APPLICANT: Cech, Thomas R.
Lingner, Joachim
Nakamura, Toru
Chapman, Karen B.
Morin, Gregg B.
Harley, Calvin B.
Handrews, William H.
TITLE OF INVENTION: Human Telomerase Catalytic Subunit
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: USAA

ZIP: 94111-3874

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: DAN FC Compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/721,456

FILING DATE: 12-NOV-1997

APPLICATION NUMBER: US/08/974,549A

FILING DATE: 19-NOV-1997

APPLICATION NUMBER: US 08/844,419

FILING DATE: 19-NOV-1997

APPLICATION NUMBER: US 08/846,017

FILING DATE: 18-APR-1997

APPLICATION NUMBER: US 08/846,017

FILING DATE: 05-APR-1997

APPLICATION NUMBER: US 08/851,843

FILING DATE: 05-MAY-1997

APPLICATION NUMBER: US 08/854,050

FILING DATE: 14-AUG-1997

APPLICATION NUMBER: US 08/912,951

FILING DATE: 14-AUG-1997

APPLICATION NUMBER: US 08/912,951

FILING DATE: 14-AUG-1997

APPLICATION NUMBER: US 08/915,503

FILING DATE: 11-AUG-1997

APPLICATION NUMBER: US 08/915,503

FILING DATE: 11-AUG-1997

APPLICATION NUMBER: US 08/915,503

FILING DATE: 11-AUG-1997

APPLICATION NUMBER: WO PCT/US97/17885

FILING DATE: 10-OCT-1997

APPLICATION NUMBER: WO WERE WO PCT/US97/17885

FILING DATE: 10-OCT-1997

APPLICATION NUMBER: WO WERE WO WERE WESTERNICE/OCKET WUMBER: 01-OCT-1997

APPLICATION NUMBER: 10-OCT-1997

APPLICATION NUMBER: 10-0CT-1997

APPLICATION NUMBER: 10-0CT-1997

APPLICATION NUMBER: 10-0CT-1997

APPLICATION NUMBER: 10-0CT-1997
                                                                           Indels
      3.6%; Score 16.4; DB 1;
94.4%; Pred. No. 1.4e+02;
iive 0; Mismatches 1;
                                                                                                                                                                                                                                                           RESULT 158
US-09-721-456-543/c
; Sequence 543, Application US/09721456
; Patent No. 6617110
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
                                                                                                                               149 CACCGTTCATTCTAGAGC 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (415) 576-0300
                                                                                                                                                                   18 CACCCTTCATTCTAGAGC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO: 543:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 18 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: San Francisco
STATE: California
3.6%
Query Match
Best Local Similarity 94.4%
Matches 17; Conservative
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Groskreutz, Debyra J.
TITLE OF INVENTION: Prohormone Convertase Transformed Cells and
Polypeptide Synthesis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                         ; NAME/KEY: -.18
; LOCATION: 1.18
; CTHER INFORMATION: /note= "antisense hTRT molecule"
; SEQUENCE DESCRIPTION: SEQ ID NO: 543:
US-09-721-456-543
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1; Indels
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3.6%; Score 16.4; DB 1;
Best Local Similarity 94.4%; Pred. No. 1.4e+02;
Matches 17; Conservative 0; Mismatches 1;
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REGISTRATION NUMBER: 34,659
REFERENCE/DOCKET NUMBER: P0748P3
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-9801
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
LENGTH: 21 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: Linear SEQ ID NO: 13: US-08-026-143B-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech, Inc.
STREET: 1 DNA Way
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 13, Application US/08026143B Patent No. 6348327 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: South San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                149 CACCGTTCATTCTAGAGC 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: DNA
PEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18 CACCCTTCATTCTAGAGC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 57
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-026-143B-13
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3.6%; Score 16.2; DB 1; Length 21; 85.7%; Pred. No. 1.6e+02; tive 0; Mismatches 3; Indels

Query Match
Best Local Similarity 85.7
Matches 18; Conservative

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Sequence 13, Application PC/TUS9210621
GENERAL INFORMATION:
APPLICANT: Genentech, Inc.
APPLICANT: Gorman, Cornelia M.,
APPLICANT: Marriott, Dave,
APPLICANT: Marriott, Dave,
APPLICANT: Groskreutz Debrya J.
TITLE OF INVENTION: Prohormone Convertase Transformed Cells and Polypeptide Synthes
NUMBER OF SEQUENCES: 54
CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech, Inc.
ADRESSEE: Genentech, Inc.
ATREET: 460 Point San Bruno Blvd
CITY: South San Francisco
STREET: Allfornia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 13, Application PC/TUS9402233
| GENERAL INFORMATION:
| APPLICANT: Genentech, Inc.
| TILE OF INVENTION: Prohormone Convertage Transformed Cells and Polypeptide Synthe:
| NUMBER OF SEQUENCES: 54
| CORRESPONDENCE ADDRESS:
| ADDRESSEE: Genentech, Inc.
| STREET: 460 Point San Bruno Blvd
| CITY: South San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: 5.25 inch, 360 Kb floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CUREVIEW.

OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: patin (Genentech)

CURRENT APPLICATION DATA:

APPLICATION NUMBER: PCT/US92/10621

FILING DATE: 19921204
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
PILING DATE: 22-MAY-1992
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
PILING DATE: 07/80331
FILING DATE: 06-DEC-1992
ATORNEY/AGENT INFORMATION:
NAME: Adler, Carolyn R.
REGISTRATION NUMBER: 32,324
REFERENCE/DOCKET NUMBER: 748P2.F
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECHONE: 415/225-2614
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  156 CATTCTAGAGCAAAAAA 176
156 CATTCTAGAGCAAACAAAAA 176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 CATTCTAGAGCAAAGAGACAA 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEX: 910/371-7168
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 21 bases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 21 bases
TYPE: NUCLEIC ACID
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ropoLoGY: linear
PCT-US92-10621-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PCT-US94-02233-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 161
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MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "peptide nucleic acid (PNA),
DESCRIPTION: where (deoxy(ribose-phosphate linkages are replaced by
DESCRIPTION: N-(2-aminoethyl)glycine units linked to nucleotide bases via
DESCRIPTION: glycine amino N through a methylenecarbonyl linker"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
APPLICANT: Shay, Jerry W.
APPLICANT: Shay, Jerry W.
APPLICANT: Piatysek, Mieczyslaw A.
APPLICANT: Piatysek, Mieczyslaw A.
APPLICANT: Corey, David R.
APPLICANT: Corey, David R.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Peptide Nucleic Acids
INTMERS OF SEQUENCES: 60
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 3.5%; Score 16; DB 1; Length 16; Best Local Similarity 100.0%; Pred. No. 1.3e+02; Matches 16; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIF: 9411-3834

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
APPLICATION NUMBER: US/09/349,532
          NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-001610US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 27:
SEQUENCE CHARACTERISTICS:
LENGTH: 16 base pairs
TYPE: nucleic acid
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/838,545
FILING DATE: 09-APR-1997
APPLICATION NUMBER: US 08/630,019
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 27, Application US/09349532
Patent No. 6294650
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TELECOMONIOTANTON INFORMATION.
TELEPRONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 27:
SEQUENCE CHARACTERISTICS:
LENGTH: 16 base pairs
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ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          53 TAACTGAGAAGGGCGT 68
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-08-838-545-27
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Batent No. 6046307

GENERAL INFORMATION:
APPLICANT: Shay, Jerry W.
APPLICANT: Wright, Woodring E.
APPLICANT: Ocey, David R.
APPLICANT: Ocey, David R.
APPLICANT: No. 6646307tcn, James C.
TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
ADDRESSE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3.6%; Score 16.2; DB 1; Length 21; 85.7%; Pred. No. 1.6e+02; tive 0; Mismatches 3; Indels
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SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/838,545
FILING DATE: 09-APR-1997
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INFORMATION:
                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: 5.25 inch, 360 Kb floppy disk COMPUTER: IBM PC compatible COMPUTER: SYSTEM: PC-DOS/MS-DOS SOFTWARE: patin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/02233
                                                                                                                                                                                                                                             FILING DATE:
CLASSIFICATION NOMBER: PCI/US94/02233
FILING DATE:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Love, Richard B.
REGISTRATION NUMBER: 34,659
REFERENCE/DOCKET NUMBER: 748P3PCT
TELECHOME: 415/25-530
TELEFAX: 415/952-9881
TELEFAX: 910/371-7168
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
LENGTH: 21 bases
TELEFAX: 12 bases
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ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      156 CATTCTAGAGCAAACAAAAA 176
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Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: single
California
: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            linear
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APPLICANT: Realey, James T.
APPLICANT: Weinrich, Scott L.
TITLE OF INVENTION: Methods for Detecting the RNA Component of TITLE OF INVENTION: Telomerase
NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
ADDRESSER: TWONSEND and TOWNSEND and CREW LLP
STREET: Two Sembarcadero Center, 8th Floor
CITY: San Francisco
STATE: California
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                                                                                                                                                                          3.4%; Score 15.2; DB 1; Length 20; B5.0%; Pred. No. 1.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 15; DB 1; Length 15; Pred. No. 1.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/770,565 FILING DATE: 20-DEC-1996 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-002300US
TELECOMNUMICATION INFORMATION:
TELEPHONE: 415-576-0300
                                                                                                                                                                                                                                          0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%; Prec. ....
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COMPUTER READABLE FORM:
CMEDIUM TYPE: Floppy disk
CMEDUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; Sequence 12, Application US/08630019A; Patent No. 6015710; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 10, Application US/08770565; Patent No. 5846723; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                 313 CTGTCAGCCGCGGGTCTCTC 332
                                                                                                                                                                                                                                                                                                                                                            CIGICIGCCACGGGTITCIC 20
       ORGANISM: Artificial Sequence
                                                                       ; OTHER INFORMATION: PCR primer US-09-596-938-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   152 CGTTCATTCTAGAGC 166
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 100. |
Matches 15, Conservative
                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Kim, Nam Woo APPLICANT: Wu, Fred
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 15 base pairs
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STRANDEDNESS: sing
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                                                                                                                                                                                                              Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-08-630-019A-12/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 166
US-08-770-565-10/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-08-770-565-10
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                                                                                                                                                                              Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: Bougueleret, Lydie
APPLICANT: Bougueleret, Lydie
TITLE OF INVENTION: A NUCLEIC ACID ENCODING A RETINOBLASTOMA BINDING PROTEIN (RBP-7)
TITLE OF INVENTION: A NUCLEIC ACID ENCODING A RETINOBLASTOMA BINDING PROTEIN (RBP-7)
TITLE OF INVENTION: UNUMBER: US/09/345,882
CURRENT APPLICATION NUMBER: US/09/345,882
CURRENT APPLICATION NUMBER: US 60/091,315
PRIOR APPLICATION NUMBER: US 60/091,315
PRIOR APPLICATION NUMBER: US 60/111,909
PRIOR FILING DATE: 1998-12-10
NUMBER OF SEQ ID NOS: 140
SOFTWARE PATENT PATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: primer_bind
| LOCATION: 1..20
| OTHER INFORMATION: upstream amplification primer for SEQ 34, SEQ 55, SEQ 35, SEQ 56
| US-09-345-882-76
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APPLICANT: Li, Xiao-Jiang
APPLICANT: Li, Xiao-Jiang
APPLICANT: Li, Shi-Una
TITLE OF INVENTION: Huntington Disease Cellular Model:
TITLE OF INVENTION: Stably Transfected FC12 Cells Expressing Mutant Huntingtin
TITLE OF INVENTION: Stably Transfected FC12 Cells Expressing Mutant Huntingtin
TITLE OF INVENTION: Stably Transfected FC12 Cells Expressing Mutant Huntingtin
TITLE OF INVENTION: Stably Transfected FC12 Cells Expressing Mutant Huntingtin
CURRENT APPLICATION NUMBER: US/09/596,938
CURRENT FILING DATE: 2000-06-19
FRIOR APPLICATION NUMBER: 60/140.018
FRIOR FILING DATE: 1999-06-18
NUMBER OF SEQ ID NOS: 15
SOFTWARE: FASTSEQ for Windows Version 3.0
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "peptide nucleic acid (PNA),
DESCRIPTION: where (deoxy(ribose-phosphate linkages are replaced by DESCRIPTION: N-(2-aminoethyl)glycine units linked to nucleotide bases via bascRIPTION: glycine amino N through a methylenecarbonyl linker"
US-09-349-532-27
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94.1%; Pred. No. 1.7e+02;
tive 0; Mismatches 1;
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// Patent No. 6355481
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                        53 TAACTGAGAAGGGCGT 68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 TAACTGAGAAGGGCGT 16
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Matches 16; Conservative
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LENGTH: 20
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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; Sequence 40, Application US/08630019A
Patent No. 6015710
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: 01.
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  42 TTGTCTAACCCTAAC 56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15 rrereradecerade 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: single
            COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  linear
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US-08-630-019A-18/C
; Sequence 18, Application US/08630019A
; Patent No. 6015710
; GENERAL INFORMATION:
; APPLICANT: Widght, Woodring E.
; APPLICANT: Piatyszek, Mieczyslaw A.
; APPLICANT: Corey, David
; APPLICANT: No. 6015710ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by
; TITLE OF INVENTION: Peptide Nucleic Acids
; VUMBER OF SEQUENCES: 46
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Townsend and Townsend and Crew LLP
APPLICANT: Shay, Jerry W.

APPLICANT: Wright, Woodring E.

APPLICANT: Placyszek, Mieczyslaw A.

APPLICANT: Orey, David James C.

APPLICANT: No. 6015710ton, James C.

TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Peptide Nucleic Acids

NUMBER OF SEQUENCES: 46

CORRESPONDENCE ADDRESS:

ADDRESSEE: Townsend and Townsend and Crew LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 3.3%; Score 15; DB 1; Length 15; Best Local Similarity 100.0%; Pred. No. 1.4e+02; Matches 15; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                 COUNTRY: CALLLULINGA

ZIP: 94111-3834

COUNTER READBLE FORM:
MEDIUW TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/NM-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/630,019A
FILING DATE: 09-JUN-1996
CLASSIFICATION: 536
ATTONEY/AGENT INPORMATION:
NAME: Storella, John REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944
REGISTRATION INFORMATION:
TELEPRAK: (415) 576-0200
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
                                                                                                                                                                                                                                       : Two Embarcadero Center, Bighth Floor
San Francisco
California
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CITY: San Francisco
STATE: California
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DESCRIPTION:
US-08-630-019A-12
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STATE:
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MOLECULE TYPE: other nucleic acid (DNA), BESCRIPTION: /desc = "peptide nucleic acid (DNA), BESCRIPTION: where (deoxy)ribose-phosphate linkages are replaced by DESCRIPTION: N-(2-aminoethyl)glycine units linked to nucleotide bases via DESCRIPTION: glycine amino nitrogen through a methylenecarbonyl linker"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 15;
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SOFTWARES PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NAMES:
FILNG DATE: 09-JUN-1996
CLASSIFICATION: 536
ATTONREY/AGENT INFORMATION:
NAME: SECRETLA JOHN R:
REGISTRATION NUMBER: 32,944
REFERENCE/POCKET NUMBER: 32,944
REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944
REJERRANICATION INFORMATION:
TELEPHONE: (415) 576-0200
INFORMATION FOR SEQ ID NO: 18:
SEQUENCE (A15) 576-0300
INFORMATION FOR SEQ ID NO: 18:
SEQUENCE CHARACTERISTICS:
LEMOTH: 15 base pairs
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3.3%; Score 15; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 1.4e+02;
Matches 15; Conservative 0; Mismatches 0;
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/desc = "peptide nucleic acid (PNA),
where (deoxy(ribose-phosphate linkages are replaced by
N-(2-aminoethy) alycine units linked to nucleotide bases via
glycine amino N through a methylenecarbonyl linker"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 2, Application US/08838545;
Sequence 2, Application US/08838545;
Patent No. 6046307
GENERAL INFORMATION:
APPLICANT: Wight, Woodring E.
APPLICANT: Orey, David R.
APPLICANT: Ocey, David R.
APPLICANT: No. 6046307ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by ITILE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 60
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                               LENGTH: 15 base pairs

TYPE: nucleic acid

TYPE: nucleic acid

TYPE: nucleic acid

TOPOLOGY: linear

MOLECULE TYPE: other nucleic acid

DESCRIPTION: /desc = "phosphorothioate (PS) nucleic acid"
                                                                                                                                                                                                                                               3.3%; Score 15; DB 1; Length 15; 100.0%; Pred. No. 1.4e+02; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OPERATION SYSTEM: FC-LUSS/MS-LUSS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/838,545
FILING DATE: 09-APR-1997
CLASSIFICATION NUMBER: US/08/838,545
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Stoorella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-001610US
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 2:
SEQUIBORE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 9411-3834
COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                           46 CTAACCCTAACTGAG 60
                                                                                                                                                                                                                                                                                                                                                                              15 CTAACCCTAACTGAG 1
TELEFAX: (415) 576-0300 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: other nucl
DESCRIPTION: /desc = "p
                                                                                                                                                                                                                                                                                              15, Conservative
                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DESCRIPTION:
DESCRIPTION:
DESCRIPTION:
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US-08-838-545-2/c
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DB 1; Length 15;

3.3%; Score 15;

Query Match

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/desc = "peptide nucleic acid (PNA),
where (deoxy(ribose-phosphate linkages are replaced by
N-(2-aminoethyl)glycine units linked to nucleotide bases via
glycine amino N through a methylenecarbonyl linker"
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                                            Gaps
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                                                                                                                                                                                                                                                                                                       APPLICANT: Shay, Jerry W.
APPLICANT: Wright, Woodring E.
APPLICANT: Piatysek, Mieczyslaw A.
APPLICANT: Piatysek, Mieczyslaw A.
APPLICANT: Pot Control of Mieczyslaw A.
APPLICANT: No. 6046307ton, James C.
TITLE OF INVENTION: Wodulation of Mammalian Telomerase by
TITLE OF INVENTION: Peptide Nucleic Acids
CORRESPONDENCE ADDRESS:
ADDRESSEE: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3.3%; Score 15; DB 1; Length 15;
100.0%; Pred. No. 1.4e+02;
tive 0; Mismatches 0; Indele
                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SUSTWARE: PEPLICALION BATA:
APPLICATION NUMBER: US/08/030,019
FILING DATE: 09-APR-1997
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTORNEY AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,94
REFERENCE/DOCKET NUMBER: 015389-001610US
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION FOR SEQ ID NO: 5:
SEQUIBACE CHARACTERISTICS:
INFORMATION FOR SEQ ID NO: 5:
                  Pred. No. 1.4e+02;
; Mismatches 0;
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100.0%; Pre-
                                                                                                                                                                                                                                             ; Sequence 5, Application US/08838545 ; Patent No. 6046307
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Best Local Similarity 100.0
Matches 15, Conservative
                                                                                    46 CTAACCCTAACTGAG
                Best Local Similarity 100 Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: San Francisco
STATE: California
COUNTRY: USA
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DESCRIPTION:
DESCRIPTION:
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DESCRIPTION:
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Sequence 2, Application US/09349532

Sequence 2, Application US/09349532

Retent No. 6294650

GENERAL INFORMATION:
APPLICANT: Shay, Jarry W.
APPLICANT: Wright, Woodring E.
APPLICANT: Piatyszek, Mieczyslaw A.
APPLICANT: Ocrey, David R.
APPLICANT: No. 6294650tcon, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 6

CORRESPONDENCE ADDRESS: 6

COR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3.3%; Score 15; DB 1; Length 15; 100.0%; Pred. No. 1.4e+02; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                 COMPUTER: IEBM PC compatible
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/030,019
FILING DATE: 09-APR-1997
CLASSIFICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATYORNEY/AGENT INPORMATION:
NAMME: Stoorella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/POCKET NUMBER: 015389-001610US
TELEPHONE: (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: USA

COMPUTER READABLE FORM:
MEDIUTY TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATE:
APPLICATION NUMBER: US/09/349,532
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "phosphorothioate (PS)
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US-08-838-545-45.
                                                                       COMPUTER: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk

"MPUTER: Floppy disk
"MPUTER: "MPUTER: PC-DOS/MS-DOS
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Best Local Similarity 100.0°
Matches 15; Conservative
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PRIOR APPLICATION DATA:
                                                                   California
: USA
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US-09-349-532-2/c
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STATE: Ca
COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ò
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            .;
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                     APPLICANT: Shay, Jerry W.
APPLICANT: Wright, Woodring E.
APPLICANT: Platyszek, Mieczyslaw A.
APPLICANT: Corey, David State C.
APPLICANT: No. 6046307con, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 45, Application US/08838545

Patent No. 6046307

GENERAL INFORMATION:
APPLICANT: Wright, Woodring E.
APPLICANT: Piatyszek, Mieczyslaw A.
APPLICANT: Corey, David M.
APPLICANT: No. 6046307ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 60

CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 15;
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                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE SUCURIOR ADDRESS:
ADDRESSE: Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STARE: California
COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/836,545
FILING DATE: 09-APR-1997
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTORNEY/GENT INFORMATION:
NAME: SLOTELIA, JOHN R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-001610US
FILEPHONE: (415) 576-0200
INFORMATION INFORMATION:
TELEPAX: (415) 576-0200
INFORMATION FOR SEQ ID NO: Z8:
LENGTH: 15 base pairs
FIRENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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3.3%; Score 15; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 1.4e+02;
Matches 15; Conservative 0; Mismatches 0;
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Two Embarcadero Center, Eighth Floor
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GENERAL INFORMATION:
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MOLECULE TYPE:
DESCRIPTION:
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US-08-838-545-28
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DESCRIPTION:
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US-08-838-545-45/c
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Gaps

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other nucleic acid / desc = "poptide nucleic acid (PNA), desc = "poptide nucleic acid (PNA), where (deoxy(ribose-phosphate linkages are replaced by N-(2-aminoethyl)glycine units linked to nucleotide bases via glycine amino N through a methylenecarbonyl linker"
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/desc = "peptide nucleic acid (PNA),
where (deoxy(ribose-phosphate linkages are replaced by
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Shay, Jerry W.
APPLICANT: Wright, Woodring E.
APPLICANT: Wright, Woodring E.
APPLICANT: Platyszek, Mieczyslaw A.
APPLICANT: Corey, David R.
APPLICANT: No. 6294650ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by
TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 60
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                  Length 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Indels
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SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/349,532
FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/838,545
FILING DATE: 09-APR-1997
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTONNEY/AGENT INFORMATION:
NAME: SECRETA, JOHN R.
REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 015389-001610US
TELEPHONE: (415) 576-0300
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 28:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                Query Match
3.3%; Score 15; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 1.4e+02;
Matches 15; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 28, Application US/09349532
Patent No. 6294650
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       42 TIGICTAACCCTAAC 56
          TELEFAX: (415) 576-0300 INFORMATION FOR SEQ ID NO: ESEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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STRANDEDNESS: single
                                                                                                                                                                         linear
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                                                                                                                                                                   TOPOLOGY: lin
MOLECULE TYPE:
DESCRIPTION:
DESCRIPTION:
DESCRIPTION:
DESCRIPTION:
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DESCRIPTION:
DESCRIPTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         other nucleic acid (PNA),

desc = "peptide nucleic acid (PNA),
where (deoxy(ribose-phosphate linkages are replaced by
N-(2-aminoethy) alycine units linked to nucleotide bases via
glycine amino N through a methylenecarbonyl linker"
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Fatent No. 6294650
GENERAL INFORMATION:
APPLICANT: Shay, Jerry W.
APPLICANT: Piatyszek, Mieczyslaw A.
APPLICANT: Octey, David R.
APPLICANT: Octey, David R.
APPLICANT: No. 6294650ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 60
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 3.3%; Score 15; DB 1; Length 15; Best Local Similarity 100.0%; Pred. No. 1.4e+02; Matches 15; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 94111-3834

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/349,532
APPLICATION NUMBER: US 08/838,545
FILING DATE: 09-APR-1997
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-001610US
TELEPANCH: (415) 576-0200
TELEPANCH: (415) 576-0200
TELEPANCH: (415) 576-0200
TELEPANCH: (415) 576-0200
TELEPANCH: 15 DATE OF ORDER
TYPE: nucleic acid
SEQUENCE CHARACTERISTICS:
LENGTH: 15 DATE DATE
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLLOGY: linear
MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "Poptide nucleic acid
CONTENTION: /desc = "Poptide nucleic acid
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CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1997
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-001610US
TELECOMMUNICATION INPORMATION:
TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        46 CTAACCCTAACTGAG 60
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DESCRIPTION: N
DESCRIPTION: G
US-09-349-532-2
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Best Local Similarity 88.9
Matches 16; Conservative
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; OTHER INFORMATION: PRIMER
US-09-673-298-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA ORGANISM: UNKNOWN
                                                    JS-09-673-298-4/c
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N-(2-aminoethyl)glycine units linked to nucleotide bases via glycine amino N through a methylenecarbonyl linker"
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                                                                                                                                                                                                                                                                                                                            Sequence 45, Application US/09349532;
Sequence 45, Application US/09349532;
Batent No. 6294650
GENERAL INFORMATION:
APPLICAMT: Wright, Woodring E.
APPLICAMT: Piatyszek, Mieczyslaw A.
APPLICAMT: Orey, David R.
APPLICAMT: No. 6294650ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 60
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                .
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 94111-3834

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION NATR:
APPLICATION NUMBER: US/09/349,532
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PEILING DATE:
CLASSIFICATION
PRIOR APPLICATION
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1997
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-001610US
TELEPHONE: (415) 576-0200
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 45:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /desc = "phosphorothioate (PS)
nucleic acid"
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j DESCRIPTION:
j DESCRIPTION:
US-09-349-532-28
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DESCRIPTION:
                                                                                                                                                                                                                                                                                                                          US-09-349-532-45/C
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other nucleic acid
/desc = "peptide nucleic acid (PNA),
where (deoxy)ribose-phosphate linkages are replaced by
N-(2-aminoethyl) glycine units linked to nucleotide bases via
glycine amino nitrogen through a methylenecarbonyl linker"
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GENERAL INFORMATION:
APPLICANT: Philip Dan Cook
TITLE OF INVENTION: PNA-DNA-PNA Chimeric
TITLE OF INVENTION: Macromolecules
NUMBER OF SEQUENCES:
ADDRESSEE: Woodcock Washburn Kurtz Mackiewicz and
ADDRESSEE: No. 5700922x1s
STREET: One Liberty Place - 46th Floor
CITY: Philadelphia
STATE: PA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 3.1%; Score 14; DB 1; Length 14; Best Local Similarity 100.0%; Pred. No. 1.6e+02; Matches 14; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATE:
APPLICATION NUMBER: US/08/630,019A
FILING DATE: 09-JUN 1996
CLASSIFICATION: 536
ATTORNEY AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 015389-001600US
TELEPAN: (415) 576-020
TELEPAN: (415) 576-020
INFORMATION FOR SEQ ID NO: 25:
SEQUENCE CHARACTERISTICS:
LENGTH: 14 base pairs
TYPE: nucleic acid
TYPE: nucleic acid
TYPE: ACCOUNT ACCO
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: Floppy disk, 1.44 Mb storage COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: WordPerfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/158,352
FILING DATE: herewith
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US92/11339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 182
US-08-158-352-2
; Sequence 2, Application US/08158352
; Patent No. 5700922
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   52 CTAACTGAGAAGGG 65
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                                                    STREET: Two bullers CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  linear
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DESCRIPTION:
DESCRIPTION:
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DESCRIPTION:
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                                                                                                                                                                      COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 3.3%; Score 14.8; DB 1; Length 19; Best Local Similarity 88.9%; Pred. No. 1.8e+02; Matches 16; Conservative 0; Mismatches 2; Indels
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APPLICANT: Shay, Jerry W.
APPLICANT: Waight, Woodring E.
APPLICANT: Parigasek, Mieczyslaw A.
APPLICANT: Posty, David
APPLICANT: No. 6015710ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by
TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Xu, Minzhen
APPLICANT: Xu, Gang
APPLICANT: Yu, Gang
APPLICANT: Humbhreys, Robert
ITILE CANT: Humbhreys, Robert
ITILE OF INVENTION: CANCER CELL VACCINE
CURRENT APPLICATION NUMBER: 03/09/205,995, (CIP)
CURRENT FILING DATE: 1998-12-04
PRIOR APPLICATION NUMBER: 09/036,746
PRIOR APPLICATION NUMBER: 09/61,627
PRIOR APPLICATION NUMBER: 09/661,627
PRIOR APPLICATION NUMBER: 1996-06-11
NUMBER OF SEQ ID NOS: 79
SEQ ID NO S: 79
SEQ ID NO S: 79
LENGTH: 18
                                                                                      TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "PRIMER"
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-08-630-019A-25/c; Sequence 25, Application US/08630019A; Patent No. 6015710
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 24, Application US/09205995; Patent No. 6368855; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           348 GTTCAGGCCTTTCAGGCC 365
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA ORGANISM: Artificial Sequence
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 19 base pairs
                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                   ORIGINAL SOURCE:
ORGANISM: Hom
                                                                                                                                                                                                                                                                                                       ANTI-SENSE: NO
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MODULATION OF GENE EXPRESSION
IN PLANTS
1263
                                                                                                                                                               CITY: Los Angeles
COUNTY: Los Angeles
STATE: California
COUNTY: U.S.A.
ZIP: 90071-2066
ZOMPUTER: READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
COMPUTER: IBW Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/679,645
FILING DATE: July 12, 1996
CLASSIFICATION NUMBER: 60/001,135
FILING DATE: July 13, 1995
APPLICATION NUMBER: 60/001,135
FILING DATE: July 13, 1995
APPLICATION NUMBER: 60/001,135
FILING DATE: September 2, 1994
ATTORNEY/AGENT INFORMATION:
REGISTRATION NUMBER: 32,327
TELECOMMUTATION: NUMBER: 32,327
TELECOMMUTATION: NUMBER: 32,327
TELECOMMUTATION: NUMBER: 32,327
TELECOMMUTATION: NUMBER: 32,327
       TITLE OF INVENTION: MODULATION OF TITLE OF INVENTION: IN PLANTS
NUMBER OF SEQUENCES: 1263
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 571, Application US/08679645
Patent No. 6350934
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity 64.7
Matches 11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      linear
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                                                                                                                                                                                                                                                                                                                                                                                                 3.1%; Score 13.8; DB 1; Length 17; 88.2%; Pred. No. 1.9e+02; tive 0; Mismatches 2; Indels
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Patent No. 6277603

GENERAL INFORMATION:

APPLICANT: Cook, Phillip Dan

FILE OF INVENTION: PNA-DNA-PNA Chimeric Macromolecules
FILE REPERENCE: ISIS3102

CURRENT APPLICATION NUMBER: US/09/108,911

CURRENT APPLICATION NUMBER: US/09/108,911

CURRENT APPLICATION NUMBER: 08/877,317

PRIOR PILING DATE: 1997-06-17

NUMBER: OF SEQ ID NOS: 4

SOFTWARE: Patentin version 3.0

LENTH: 17
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3.1%; Score 13.8; DB 1; Length 17;
Best Local Similarity 88.2%; Pred. No. 1.9e+02;
Matches 15; Conservative 0; Mismatches 2; Indels
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APPLICANT: Edington, Brent E.
APPLICANT: McSwiggton, James A.
APPLICANT: McSwiggton, James An Owens
APPLICANT: Merlo, Patricia Ann Owens
APPLICANT: Skokut, Thomas A.
APPLICANT: Skokut, Thomas A.
APPLICANT: Folkerts, Otto
APPLICANT: Folkerts, Otto
APPLICANT: Merlo, Donald J.
TITLE OF INVENTION: COMPOSITION AND METHODS FOR
                                                                                                  ISIS-1236
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; NAME/KEY: misc_feature
; OTHER INFORMATION: Antisense Sequence
US-09-108-911-2
FILING DATE: December 23, 1992
ATTORNEY/AGENT INFORMATION:
NAME: JOHN W. Caldwell
REGISTRATION NUMBER: 28, 937
REFERENCE/DOCKET NUMBER: 151S-12:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 215-568-3100
TELEPHONE: 215-568-349
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 184
US-08-679-645-826
; Sequence 826, Application US/08679645
; Patent No. 6350934
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          102 TTCTCGCTGACTTTCAG 118
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                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 88.2*
...rhes 15; Conservative
                                                                                                                                                                                                                                                          TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
ANTI-SENSE: yes
                                                                                                                                                                                                                                                                                                                                                      US-08-158-352-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-09-108-911-2
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Gaps
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0
3.1%; Score 13.8; DB 1; Length 17; 64.7%; Pred. No. 1.9e+02; ive 4; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                           merlo, Patricia Ann Owens

Jach, James A.

Merlo, Inling
APPLICANT: Skokut, Thomas A.
APPLICANT: Skokut, Thomas A.
APPLICANT: Folkerts, Otto
APPLICANT: Merlo, Donald J.
TITLE OF INVENTION: COMPOSITION AND METHODS FOR
TITLE OF INVENTION: MODULATION OF GENE EXPRESSION
TITLE OF INVENTION: IN PLANTS
TITLE OF INVENTION: IN PLANTS
MUMBER OF SEQUENCES: 1263
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West F.
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US-10-232-634-11
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; Sequence 11, Application US/09586376
; Patent No. 6492115
; GENERAL INFORMATION:
; APPLICANT: Guida, Marco
; APPLICANT: Hall, Jeff
; TITLE OF INVENTION: AND RELATED MATERIALS AND METHODS
; FILE REFERENCE: 4389-20
; CURRENT FAPLICATION UNMBER: US/09/586,376
; CURRENT FILING DATE: 2000-06-02
; WUMBER OF SEQ ID NOS: 29
; SOFTWARE: Patentin Ver. 2.1
; SEQ ID NO 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
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Pred. No. 2e+02;
0; Mismatches 1; Indels
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Mismatches
                                                            COUNTRY: U.S.A.

ZIP: 90071-206
COMPUTER READABLE FORM:
MEDIUM TYPE: STORAGE
COMPUTER: 13.5" Diskette, 1.44 Mb
MEDIUM TYPE: STORAGE
COMPUTER: 18 Compatible
OPERATING SYSTEM: 1BM P.C. DOS S.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/679,645
FILING DATE: July 12, 1996
CLASSIFICATION DATA:
APPLICATION NUMBER: 08/100,726
FILING DATE: SEPTEMBER: 2, 1994
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 219/247
TELEPHONE: (213) 489-1600
TELEFAX: (213) 489-1600
TELEFAX: (213) 955-0440
TELE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
3.1%; Score 13.8;
Best Local Similarity 82.4%; Pred. No. 2e
Matches 14; Conservative 1; Mismatche
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Best Local Similarity 93.3%;
Matches 14; Conservative
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Los Angeles
California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-09-586-376-11
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Sequence 4559, Application US/09371772B
; Sequence 4559, Application US/09371772B
; Sequence 4559, Application US/09371772B
; Sequence 4559, Application
; Patent No. 656127:
; GENERAL INFORMATION:
; APPLICANT: Parco, Pan
; APPLICANT: Parco, Pan
; APPLICANT: Stinchcomb, Dan
; APPLICANT: Stinchcomb, Dan
; APPLICANT: Stinchcomb, Dan
; APPLICANT: Stinchcomb, Dan
; TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re
; TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
; TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
; TITLE OF INVENTION: 1999-08-10
; FILE REFERENCE: 1999-08-10
; FILE REFERENCE: 1995-10-26
; PRIOR FILING DATE: 1995-10-26
; PRIOR FILING DATE: 1996-01-08
; NUMBER OF SEQ ID NOS: 14225
; SOFTWARE: PatentIn version 3.0
; SEQ ID NO 4559
: LENGTH: 17
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Sequence 12, Application US/09586376
Patent No. 6492115
GENERAL INFORMATION
GENERAL INFORMATION
APPLICANT: Hall, Jeff
TITLE OF INVENTION: GENETIC TYPING OF THE HUMAN CYTOCHROME.P450 2A6 GENE
TITLE OF INVENTION: AND RELATED MATERIALS AND METHODS
FILE REFERENCE: 4389-20
CURRENT FILING DATE: 2000-06-02
NUMBER OF SEQ ID NOS: 29
SOFTWARE: Patentin Ver. 2.1
LENGTH: 17
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Patent No. 6797477

GENERAL INFORMATION:

APPLICANT: Guida, Marco

APPLICANT: Hall, Jeff

TITLE OF INVENTION: GENETIC TYPING OF THE HUMAN CYTOCHROME P450 2A6 GENE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gape
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3.0%; Score 13.4; DB 1; Length 17; 93.3%; Pred. No. 2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Mismatches
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Query Match
Best Local Similarity 93.3
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                     ; ORGANISM: Homo sapiens
US-09-586-376-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; ORGANISM: Homo sapiens
US-09-371-772B-4559
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46 CTAACCCTAACTG 58
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy of
                                                                                                                                                                                                                                                                                                                                                                                  linear
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Best Local Similarity
Matches 13; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-08-630-019A-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY:
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APPLICANT: Guida, Marco
APPLICANT: Guida, Marco
APPLICANT: Guida, Marco
APPLICANT: Guida, Marco
APPLICANT: Hall, Jeff
TITLE OF INVENTION: GENETIC TYPING OF THE HUMAN CYTOCHROME P450 2A6 GENE
TITLE OF INVENTION: AND RELATED MATERIALS AND METHODS
TITLE REPERENCE: 4389-20
CURRENT APPLICATION NUMBER: US/10/232,634
CURRENT FILING DATE: 2002-06-02
PRIOR APPLICATION NUMBER: US/09/586,376
NUMBER OF SEQ ID NOS: 29
SOFTWARE: Patentin Ver: 2.1
SEQ ID NOS: 29
                                                                                                                                                                                                                                                                            Gaps
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3.0%; Score 13.4; DB 1; Length 17;
Best Local Similarity 93.3%; Pred. No. 2e+02;
Matches 14; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3.0%; Score 13.4; DB 1; Length 17; 93.3%; Pred. No. 2e+02; tive 0; Mismatches 1; Indels
TITLE OF INVENTION: AND RELATED MATERIALS AND METHODS FILE REPERENCE: 4389-20
CURRENT APPLICATION: 4389-20
CURRENT FILING DATE: 2002-08-30
PRIOR APPLICATION NUMBER: US/09/586,376
PRIOR FILING DATE: 2000-06-02
NUMBER OF SEQ ID NOS: 29
SOFTWARE: Patentin Ver: 2.1
LENGTH: 17
                                                                                                                                                                                                                                                                                                                                                                                  RESULT 190
US-10-232-634-12
Sequence 12, Application US/10232634
Patent No. 6797477
                                                                                                                                                                                                                                                                                                    41 TITGICTAACCCTAA 55
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                                                                                                                                                                              ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
Matches 14; Conserv
                                                                                                                                                                                                   US-10-232-634-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; UKGANISM: DOI
US-10-232-634-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY:
                                                                                                                                                                    TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 191
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Length 13;
          DB 1; Lens.
1. 1.7e+02;
0;
                                                                    ilarity 100.0%; Pred. No. 1.7
Conservative 0; Mismatches
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                                     2.9%; Score 13;
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Wright, Woodring E.
Piatyszek, Mieczyslaw A.
Corey, David R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-888-545-1/c
; Sequence 1, Application US/08838545
; Patent No. 6046307
; GENERAL INFORMATION:
; APPLICANT: Shay, Jerry W.
; APPLICANT: Wright, Woodring E.
; APPLICANT: Piatyszek, Mieczyslaw A
; APPLICANT: Corey, David R.
                                                                                                                                                                                                                                                                                                                                                                   US-08-630-019A-41/c
; Sequence 41, Application US/08630019A
Patent No. 6015710
; GENERAL INFORMATION:
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Best Local Similarity 100.
Matches 13; Conservative
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      Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                  other nucleic acid (GNA),
/desc = "peptide nucleic acid (FNA),
where (deoxy)ribose-phosphate linkages are replaced by
N-(2-aminoethyl) alycine units linked to nucleotide bases via
glycine amino nitrogen through a methylenecarbonyl linker"
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/desc = "peptide nucleic acid (PNA),

where (descy)ribose-phosphate linkages are replaced by

N.(2-aminoethyl)sylvcine units linked to nucleotide bases via

glycine amino nitrogen through a methylenecarbonyl linker"
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Sequence 17, Application US/08630019A

Patent No. 6015710X:
APPLICANT: Shay, Jerry W.
APPLICANT: Patrysek, Mieczyslaw A.
APPLICANT: Porcy, David
APPLICANT: Octey, David
APPLICANT: Corey, David
APPLICANT: Corey, David
APPLICANT: Corey, David
APPLICANT: Corey, David
APPLICANT: Wo. 6015710ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by
TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 46
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STREET: Correct
STREET: C
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MEDIUM TYPE: Floppy disk

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-COMPATIBLE

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/630,019A

FILING DATE: 09-JUN-1996

CLASSIFICATION: 536

ATTORNEY AGENT INFORMATION:

NAME: Storella, John R.

REGISTRATION NUMBER: 015389-001600US

TELEPHONE: (415) 576-0300

TELEPHONE: (415) 576-0300

INFORMATION FOR SEQ ID NO: 17:

SEQUENCE CHARACTERISTICS:

LINFORMATION FOR SEQ ID NO: 17:

SEGUENCE CHARACTERISTICS:

LEMGTH: 13 base pairs

TYPE: muclalc acid

STRANDEDNESS: single
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TELEPAX: (415) 576-0200
TELEPAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
LENGTH: 13 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc - "---
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DESCRIPTION:
DESCRIPTION:
DESCRIPTION:
DESCRIPTION:
US-08-630-019A-17
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DESCRIPTION:
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STATE: Ca
COUNTRY:
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                                                                                                                                                                                                                                                                                                                                               APPLICANT: Shay, Jerry W.
APPLICANT: Wishel, Woodring E.
APPLICANT: Platy Medging E.
APPLICANT: Platy Sek, Micczyslaw A.
APPLICANT: Platy Sex Addition of Mammalian Telomerase by TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Peptide Nucleic Acids CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco STATE: California COUNTRY: USA
ZIP: 94111-3834
ZIP: 9411-3834
ZIP: Ploppy disk.
MEDIUM TYPE: Floppy disk.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear | TOPOLOGY: linear | TOPOLOGY: linear | MOLECULE TYPE: other nucleic acid | DESCRIPTION: /desc = "phosphorothioate (PS) nucleic acid" US-08-630-019A-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2.9%; Score 13; DB 1; Length 13; 100.0%; Pred. No. 1.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: F.LOPPY GLEAR

COMPTURE: IBM PC Compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOCTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/630,019A

FILING DATE: 09-JUN-1996

CLASSIFICATION: 536

ATTORNEY/AGBNT INFORMATION:

NAMME: Storella, John R.

REGISTRATION NUMBER: 015389-001600US

TELEPHONE: 756-0200

TELEPHONE: (415) 576-0300

INFORMATION FOR SEQ ID NO: 41:

SEQUENCE CHARACTERISTICS:

LENGTH: 13 base pairs

TYPE: nucleic acid

STRANDEDNESS: single
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Query Match
2.9%; Score 13; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 1.7e+02;
Matches 13; Conservative 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 13 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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MOLECULE TYPE:
DESCRIPTION:
DESCRIPTION:
DESCRIPTION:
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where (deoxy(ribose-phosphate linkages are replaced by
N-(2-aminoethyl) glycine units linked to nucleotide bases via
glycine amino N through a methylenecarbonyl linker"
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APPLICANT: No. 6046307ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 60
CORRESPONDENCES: BOURDESS: ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco Conter, Eighth Floor CITY: San STATE: California
COUNTY: USA
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APPLICANT: Shay, Jerry W.
APPLICANT: Muright, Woodring E.
APPLICANT: Piatyszek, Mieczyslaw A.
APPLICANT: Piatyszek, Mieczyslaw A.
APPLICANT: Corey, David R.
APPLICANT: Corey, David R.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 60
CORRESPONDENCES. 60
CORRESPONDENCES. 100
STREET: Two members and Townsend and Crew LLP
STREET: Two Embaccadero Center, Eighth Floor
CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 13;
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                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/838,545
FILING DATE: 09-APR-1997
CLEASIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INFORMATION:
NAME: SLOCELIA, John R.
REGISTRATION NUMBER: 32,944
REPREMENCE/DOCKET NUMBER: 015389-001610US
TELEPHONE: (415) 576-0200
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 13 base pairs
TYPE: MICHAEL CATABLE
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2.9%; Score 13; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 1.7e+02;
Matches 13; Conservative 0; Mismatches 0;
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COMPUTER READABLE FORM:
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MOLECULE TYPE:
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US-08-838-545-4/c
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/desc = "peptide nucleic acid (PNB),
where (deoxy/ribose-phosphate linkages are replaced by
N-(2-aminoethyl)glycine units linked to nucleotide bases via
glycine amino N through a methylenecarbonyl linker"
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APPLICANT: Shay, Jerry W.
APPLICANT: Wright, Woodring E.
APPLICANT: Piatyszek, Mieczyslaw A.
APPLICANT: Corey, David R.
APPLICANT: Corey, David R.
APPLICANT: No. 6046307ton, James C.
TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 60
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 13;
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MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/838,545

FLING DATE: 09-APR.1997

CLASSIFICATION 536

PRIOR APPLICATION DATE: US/08/838,545

PILING DATE: 09-APR.1997

APPLICATION NUMBER: US/08/630,019

FILING DATE: 09-APR.1996

ATOMNEY/AGENT INPORMATION:

NAME: Storella, John R.

REGISTRATION NUMBER: 32,944

REGISTRATION NUMBER: 32,944

REGISTRATION NUMBER: 35,046

TELEPAK: (415) 576-0300

INFORMATION FOR SEQ ID NO: 4:

SEQUIBLEE CHARATICHS: 12,0000

INFORMATION FOR SEQ ID NO: 4:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 09-APR-1997
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INFORMATION:
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                                            ; DESCRIPTION:
; DESCRIPTION:
US-08-838-545-46
            MOLECULE TYPE:
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                                                                                                                                                                                                           Query Match
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where (decxy/ribose-phosphate linkages are replaced by
N-(2-aminoethyl) glycine units linked to nucleotide bases via
glycine amino N through a methylenecarbonyl linker"
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CORRESPONDENCE ADDRESS: ADDRESSE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2.9%; Score 13; DB 1; Length 13; 100.0%; Pred. No. 1.7e+02; Live 0; Mismatches 0; Indels
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ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/838,545
FILING DATE: 09-APR-1997
CLASSIFICATION 536
PRIOR APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTONINY/AGENT INPORMATION:
NAME: AFACALLE AFACALLE AFACALLE ATTONING AT
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-001610US
TELECOMMUNICATION INPORMATION:
TELEPAX: (415) 576-0300
TELEPAX: (415) 576-0300
INPORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
LENGTH: 13 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPDLOGY: linear
MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "peptide nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ALICKNES FACETIA, JOHN R.

REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944
RECISTRATION NUMBER: 015389-001610US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 46:
SEQUENCE CHARACTERISTICS:
LENGTH: 13 base pairs
TYPE: nucleic acid
STRANDENNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 198
US-08-838-545-46/c
; Sequence 46, Application US/08838545
; Patent No. 6046307
; GENERAL INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   44 GICTAACCCIAAC 56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GTCTAACCCTAAC 1
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Matches 13; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DESCRIPTION:
DESCRIPTION:
DESCRIPTION:
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/desc = "peptide nucleic acid (PNA), where (deoxy/ribose-phosphate linkages are replaced by N-(2-aminoethyl)glycine units linked to nucleotide bases via glycine amino N through a methylenecarbonyl linker"
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                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Shay, Jerry W.
APPLICANT: Wight, Woodring E.
APPLICANT: Piarysek, Mieczyslaw A.
APPLICANT: Corey, David R.
APPLICANT: Corey, David R.
APPLICANT: No. 6046307ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by NUMBER OF SEQUENCES: 60
CORRESPONDENCE ADDRESS:
                                                                                                     DB 1; Length 13;
1.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2.9%; Score 13; DB 1; Length 13; 100.0%; Pred. No. 1.7e+02; Live 0; Mismatches 0; Indels
                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CAPPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
COMPUTER: EN PC Compatible
OPERATIONS SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/838,545
FILING DATE: 09-APR-1997
CLASSIFICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INFORMATION:
NAME: SCOTELLA, JOHN R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-001610US
TELEPAN: (415) 576-0200
TELEPAN: (415) 576-0300
TELEPAN: CRARACTERISTICS:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
               /desc = "phosphorothioate (PS) nucleic acid"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Crew
                                                                                                                                                0; Mismatches
                                                                                                   2.9%; Score 13;
100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Townsend and Townsend and
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other nucleic acid
                                                                                                                                                                                                                                                                                                                                           Sequence 51, Application US/08838545
Patent No. 6046307
GENERAL INFORMATION:
                                                                                                                         100.08;
                                                                                                                                                                                           42 TTGTCTAACCCTA 54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   38 TITITIGICIAAC 50
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                                                                                                                                                13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 13 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                 13 rrercraaccera 1
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Best Local Similarity
                                                                                                                            Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  94111-3834
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DESCRIPTION:
DESCRIPTION:
DESCRIPTION:
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COMPUTER READABLE FORM:
COMPUTER READABLE FORM:
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      53 TAACTGAGAAGGG 65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
les 13; Conserv
                                                                                        STATE: Ca
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY:
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         other nucleic acid (PNA),
/desc = "peptide nucleic acid (PNA),
where (deoxy(ribose-phosphate linkages are replaced by
N-(2-aminoethy))glycine units linked to nucleotide bases via
glycine amino N through a methylenecarbonyl linker"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   .;
0
                           Sequence 52, Application US/08838545
Patent No. 6046307
GENERAL INPORMATION:
APPLICANT: Shay, Jerry W.
APPLICANT: Corey, David R.
APPLICANT: Corey, David R.
APPLICANT: Corey, David R.
APPLICANT: No. 6046307ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 60
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2.9%; Score 13; DB 1; Length 13; 100.0%; Pred. No. 1.7e+02; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                              OFFMANING SYSIEM:
CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/838,545
FILING DATE: 09-APR-1997
FILING DATE: 09-APR-1997
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTORNEY/AGBNT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-001610US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 52:
SEQUIBACE CHARACTERICATICS:
                                                                                                                                                                                                                                                              ADDRESSEE: Townsend and Townsend and Crew LL STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Shay, Jerry W.

Wright, Woodring E.

Patyszek, Mieczyslaw A.

Corey, David R.

No. 6046307ton, James C.
                                                                                                                                                                                                                                                                                                                    STATE: California
CONTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: TIM PC compatible
OFFERTING SYSTEM: PC-DOS/MS-DOS
OFFERTING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                39 TTTTTGTCTAACC 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 13 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           13 TTTTGTCTAACC 1
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DESCRIPTION:
DESCRIPTION:
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DESCRIPTION:
RESULT 200
US-08-838-545-52/c
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TOPOLOGY: linear
MOLBCULE TYPE: other nucleic acid (PNA),
DESCRIPTION: /desc = "peptide nucleic acid (PNA),
DESCRIPTION: where (decoxy(ribose-phosphate linkages are replaced by
DESCRIPTION: where (decoxy(ribose-phosphate linkages are replaced by
DESCRIPTION: N-(2-aminoethyl)glycine units linked to nucleotide bases via
DESCRIPTION: glycine amino N through a methylenecarbonyl linker"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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NUMBER OF SEQUENCES: 60
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Bighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 13;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/838,545
FILING DATE: 09-APR-1997
CLASSIFICATION: 536
PRIOR APPLICATION S36
PILING DATE: 09-APR-1996
ATTONNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REGISTRATION STOCKET NUMBER: 015389-001610US
TELEPAN: (415) 576-0200
TELEPAN: (415) 576-0300
INFORMATION FOR SEQ ID NO: 55:
SEQUENCE CHARACTERISTICS:
LENGTH: 13 base pairs
TURED THE DATE DATE: US A STOREL STO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2.9%; Score 13; DB 1; Le
100.0%; Pred. No. 1.7e+02;
tive 0; Mismatches 0;
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/desc = "peptide nucleic acid (PNB), where (decxy/tibose-phosphate linkages are replaced by N-(2-aminoethyl)glycine units linked to nucleotide bases via glycine amino N through a methylenecarbonyl linker"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2.9%; Score 13; DB 1; Le
100.0%; Pred. No. 1.7e+02;
iive 0; Mismatches 0;
                                 NAME: Scorella, John R.
REGISTATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-001610US
TELEFONMUNICATION INPORMATION:
TELEFONMUNICATION 1NPORMATION:
TELEFAX: (415) 576-0300
INPORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 13 base pairs
TYPE: nucleic acid
STRANDENNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/838,545
FILING DATE: 09-APR-1997
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                          other nucleic acid
            ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity 100.
1es 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     46 CTAACCCTAACTG 58
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                                                                                                                                                                                                                                                                                                                                                                  linear
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DESCRIPTION:
DESCRIPTION:
DESCRIPTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-09-349-532-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 other nucleic acid
/desc = "peptide nucleic acid (PNA),
where (deoxy(ribose-phosphate linkages are replaced by
N-(2-aminoethy) alycine units linked to nucleotide bases via
glycine amino N through a methylenecarbonyl linker"
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US-09-349-532-1/c

Sequence 1, Application US/09349532

Sequence 1, Application US/09349532

GENERAL INFORMATION:
APPLICANT: Shay, Jerry W.
APPLICANT: Piatyszek, Mieczyelaw A.
APPLICANT: Corey, David R.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
COMPUTRY: USA

ZIP: 94111-3834

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
COMPUTER PEADABLE FORM:
COMPUTER P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 2.9%; Score 13; DB 1; Length 13; Best Local Similarity 100.0%; Pred. No. 1.7e+02; Matches 13; Conservative 0; Mismatches 0; Indels
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COBERATING SYSTEM: PC-DOS/MS-DOS
SOCTAMER PETENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/838,545
FILING DATE: 09-APR-1997
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/630,019
FILING DATE: 00-APR-1996
ATORNEY AGENT INFORMATION:
NAME: Storella, John R.
REFERENCE/DOCKET NUMBER: 015389-001610US
TELEPHONE: (415) 576-0300
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 56:
SEQUENCE CHARACTERISTICS:
LENGTH: 13 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/09/349,532
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE:
CLASSIFICATION:
PLASSIFICATION DATA:
APPLICATION NUMBER: US 08/838,545
FILING DATE: 09-APR-1997
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  55 ACTGAGAAGGGCG 67
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: 11.
MOLECULE TYPE:
DESCRIPTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DESCRIPTION:
DESCRIPTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DESCRIPTION:
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Floppy disk
              Query Match
Best Local Similarity 100.(
Matches 13, Conservative
                                                                                                                  44 GICTAACCCIAAC 56
                                                                                                                                                               GTCTAACCCTAAC 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 13 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                linear
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Matches 13; Conserv
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                                                                                                                                                       d
TYPE: nuccompages: single
STRANDENESS: single
TOPOLOGY: linear
MOLECULE TYPE: other nucleic acid (PNA),
DESCRIPTION: /desc = "peptide nucleic acid (PNA),
DESCRIPTION: where (deoxy/ribose-phosphate linkeds are replaced by
DESCRIPTION: where the properties are replaced by
DESCRIPTION: where the properties are replaced by
DESCRIPTION: y-(2-aminoethy) glycine units linked to nucleotide bases via
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where (deoxy/tibose-phosphate linkages are replaced by
N-(2-aminoethyl)glycine units linked to nucleotide bases via
Glycine amino N through a methylenecarbonyl linker"
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                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Shay, Jerry W.
APPLICANT: Shay, Jerry W.
APPLICANT: Piatyszek, Mieczyslaw A.
APPLICANT: Piatyszek, Mieczyslaw A.
APPLICANT: Corey, David R.
APPLICANT: No. 6294650ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 60
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                       Query Match 2.9%; Score 13; DB 1; Length 13; Best Local Similarity 100.0%; Pred. No. 1.7e+02; Matches 13; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 94111-3834

COMPUTER READABLE FORM:
MEDIUM TYER: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30

CURREMY APPLICATION DATA:
APPLICATION NUMBER: US/09/349,532
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE:

CLASSIFICATION:

PLING DATE:

CLASSIFICATION DATA:

APPLICATION NUMBER: US 08/838,545

FILING DATE: 09-APR-1997

APPLICATION NUMBER: US 08/630,019

FILING DATE: 09-APR-1996

ATTORNEY/AGENT INFORMATION:

NAME: Storella, John R.

REGISTRATION NUMBER: 32,944

REFERENCE/DOCKET NUMBER: 32,944

TELECOMMUICATION INFORMATION:

TELEPHONE: (415) 576-0200

TELEPHONE: (415) 576-0300

INPORMATION FOR EGO ID NO: 12:

SEQUENCE CHARACTERISTICS:

SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 12, Application US/09349532
Patent No. 6294650
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        other nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                 13 TIGICTAACCCTA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      linear
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MOLECULE TYPE:
DESCRIPTION:
DESCRIPTION:
DESCRIPTION:
DESCRIPTION:
DESCRIPTION:
US-09-349-532-12
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                                                                    Gaps
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                                                                    ;
0
                                                                                                                                                                                                                                                                                                                                              APPLICANT: Shay, Jerry W.
APPLICANT: Shay, Jerry W.
APPLICANT: Shay, Jerry W.
APPLICANT: Platyszek, Mieczyslaw A.
APPLICANT: Corey, David R.
APPLICANT: David R.
APPLICANT: No. 6294650ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by
TITLE OF INVENTION: Peptide Nucleic Acids
ORRESPONDENCES: 60
CORRESPONDENCES. 60
CORRESPONDENCE ADDRESS:
ADDRESSER: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           .
0
                      Length 13;
                                                                  Indels
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                 DB 1; Le
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/349,532
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE:
PRIOR APPLICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/838,545
FILING DATE: 09-ABR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTATION NUMBER: 32,94
RECESTATION NUMBER: 32,94
REFERENCE/DOCKET NUMBER: 015389-001610US
TELECOMMUNICATION INFORMATION:
TELEFAX: (415) 576-0200
2.9%; Scor.
100.0%; Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "phosphorothioate (PS)
DESCRIPTION: nucleic acid"
                                                                                                                                                                                                                                                                              Sequence 46, Application US/09349532; Patent No. 6294650; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-09-349-532-51/c; Sequence 51, Application US/09349532
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MOLECULE TYPE: other nucleic acid DESCRIPTION: /desc = "peptide nucleic acid (PNA), DESCRIPTION: where (deoxy/ribose-phosphate linkages are replaced by DESCRIPTION: where (deoxy/ribose-phosphate linkages are replaced by DESCRIPTION: N-(2-aminoethyl) alycine units linked to nucleotide bases via DESCRIPTION: glycine amino N through a methylenecarbonyl linker"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
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APPLICANT: Wright, Woodring E.
APPLICANT: Platyszek, Mieczysław A.
APPLICANT: Corey, David R.
APPLICANT: Corey, David R.
APPLICANT: No. 6294650ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 60
CORRESPONDENCE ADDRESS:
ADDRESSER: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2.9%; Score 13; DB 1; Length 13; 100.0%; Pred. No. 1.7e+02; ive 0; Mismatches 0; Indels
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
COUNTRY: USA
ZIP: 9411-13834

COMPUTER: USA
ZIP: 9411-13834

COMPUTER: PALDABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/09/349,532
FILING DATE:
CLASSIFICATION NUMBER: US 08/838,545
FILING DATE: 09-APR-1997
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Storella John R.
REPERENCE/DOCKET NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 32,5600
TELEPHONE: (415) 576-0300
TELEPHONE: (415) 576-0300
TELEPHONE: HIS BASE PAIRE
TELEPHONE CHARACTERISTICS:
LYDRORMATION FOR SEQ ID NO: 52:
SEQUENCE CHARACTERISTICS:
LYDRORMATION LYDRORMATION LYDRORMATION LYDRORMATION LYDRORMATION LYDRORMATION LY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            IBM PC compatible
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   39 TITITGICIAACC 51
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Matches 13; Conservative
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-09-349-532-55/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
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TOPOLOGY:
TOPOLOGY:
MOLECULE TYPE: other nucleic acid (PNA),
DESCRIPTION: /desc = "peptide nucleic acid (PNA),
DESCRIPTION: where (deoxy(ribose-phosphate linkages are replaced by
DESCRIPTION: where (deoxy(ribose-phosphate linkages are replaced by
DESCRIPTION: N-(2-aminoethyl)glycine units linked to nucleotide bases via
DESCRIPTION: glycine amino N through a methylenecarbonyl linker"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
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0
                                                                                              APPLICANT: Shay, Jerry W. APPLICANT: Wright, Woodring E. APPLICANT: Wright, Woodring E. APPLICANT: Piatyszek, Mieczyslaw A. APPLICANT: Corey, David R. APPLICANT: Or. 6294650ton, James C. TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Peptide Nucleic Acids NUMBER OF SEQUENCES: 60 CORRESPONDENCE ADDRESS: ADDRESSE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-09-349-512-52/c

Sequence 52, Application US/09349532

Sequence 52, Application US/09349532

GENERAL INFORMATION:
APPLICANT: Shay, Jerry W.
APPLICANT: Wright, Woodring E.
APPLICANT: Piatyszek, Mieczyslaw A.
APPLICANT: Piatyszek, Mieczyslaw A.
APPLICANT: Orey, David R.
APPLICANT: No. 6294650ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 2.9%; Score 13; DB 1; Length 13; Best Local Similarity 100.0%; Pred. No. 1.7e+02; Matches 13; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/349,532
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIPICATION:
PRIOR APPLICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/636,545
FILING DATE: 09-APR-1997
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REFERRNCE/DOCKET NUMBER: 32,944
REFERRNCE/DOCKET NUMBER: 015389-001610US
TELECHONICATION INFORMATION:
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 51:
SEQUENCE CHARACTERISTICS:
LENGTH: 13 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         38 TTTTTGTCTAAC 50
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                                 Patent No. 6294650
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-09-349-532-51
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2.9%; Score 13; DB 1; Le
100.0%; Pred. No. 1.7e+02;
iive 0; Mismatches 0;
FILING DATE: 09-APR-1996

ATTONNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-001610US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0300
TELEPAX: (415) 576-0300
INPORMATION FOR SEQ ID NO: 56:
SEQUENCE CHARACTERISTICS:
LENGTH: 13 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: other nucleic acid DESCRIPTION: /desc = "peptide nu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 2, Application US/09657445A Patent No. 6608036
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; Sequence 8, Application US/09657445A
; Patent No. 6608036
; GENERAL INFORMATION:
; APPLICANT: Geron Corporation
; APPLICANT: Gryaznov, Sergei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 42 TIGICIAACCCIA 54
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nes 13; Conservative
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Matches 13; Conserv
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DESCRIPTION:
DESCRIPTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 211
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Matches
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where (deoxy/tibose-phosphate linkages are replaced by
N.(2-aminoethyl) alycine units linked to nucleotide bases via
glycine amino N through a methylenecarbonyl linker"
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Patent No. 6294650

GENERAL INFORMATION:
APPLICANT: Shay, Jarry W.
APPLICANT: Wight, Woodring E.
APPLICANT: Ocrey, David R.
APPLICANT: No. 6294650ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 60
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 2.9%; Score 13; DB 1; Length 13; Best Local Similarity 100.0%; Pred. No. 1.7e+02; Matches 13; Conservative 0; Mismatches 0; Indels
                               PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTY: USA
ZIP: 94111-3834
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/WS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco
                   CURRENT APPLICATION DATA:

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/349,532
FILING DATE:
CLASSIFICATION DATA:

PRIOR APPLICATION DATA:

PRIOR APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1997
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.

REGISTRATION NUMBER: 015389-001610US
TELEFRENCE/DOCKET NUMBER: 015389-001610US
TELEFAX: (415) 576-0300
TELEFAX: (415) 576-0300
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 55:
SEQUENCE CHARACTERISTICS:
LENGTH: 13 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/838,545
FILIND DATE: 09-APR-1997
APPLICATION NUMBER: US 08/630,019
       PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        other nucleic acid
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       OPERATING SYSTEM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE:
DESCRIPTION:
DESCRIPTION:
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DESCRIPTION:
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US-09-349-532-56/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY:
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GENERAL INFORMATION:
APPLICANT: Geron Corporation
APPLICANT: Geron Corporation
APPLICANT: Gryaznov, Sergei
APPLICANT: Pongracz, Krisztina
APPLICANT: Matray, Tracey
TITLE OF INVENTION: Oligonuclectide N3'-P5' Thiophosphoramidates: Their Synthesis and
TITLE OF INVENTION: Oligonuclectide N3'-P5' Thiophosphoramidates: Their Synthesis and
TITLE OF INVENTION: Oligonuclectide N3'-P5' Thiophosphoramidates: Their Synthesis and
CURRENT APPLICATION NUMBER: US 60/157,445A
PRIOR PAPLICATION NUMBER: US 60/153,201
PRIOR PAPLICATION NUMBER: US 60/160,444
PRIOR APPLICATION NUMBER: US 60/160,444
PRIOR APPLICATION NUMBER: US 60/160,444
SRIDE OF SEQ ID NOS: 9
SOFTWARE: Patentin version 3.1
SEQ ID NOS: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; OTHER INFORMATION: Synthetic oligonucleotide with potential inhibition activity US-09-657-445A-2
/desc = "peptide nucleic acid (PNA),
where (deoxy/tibose-phosphate linkages are replaced by
N-(2-aminoethyl)glycine units linked to nucleotide bases via
glycine amino N through a methylenecarbonyl linker"
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APPLICANT: FOURTH ALEXALING
APPLICANT: FOURTH ALEXALING
APPLICANT: Mattay, Tracey
TITLE OF INVENTION: Oligonucleotide N3'-->PS' Thiophosphoramidates: Their Synthesis ar
FILE REPERENCE: 039/004.

CURRENT APPLICATION NUMBER: US/10/463,076

CURRENT FILING DATE: 2003-06-17

PRIOR FILING DATE: 2000-09-08

PRIOR FILING DATE: 1999-09-10

PRIOR FILING DATE: 1999-09-10

PRIOR FILING DATE: 1999-10-19

NUMBER OF SEQ ID NOS: 9

SOFTWARE: PATENTIAN OF SEQ ID NOS: 9

SOFTWARE: PATENTIAN OF SEQ ID NOS: 9

LENGTH: 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: MENG, XING-JIN
APPLICANT: HALBUR, PATRICK G.
APPLICANT: HALBUR, PATRICK G.
APPLICANT: MOROZOV, IGOR
APPLICANT: MOROZOV, IGOR
APPLICANT: LUM, MELISSA A.
TITLE OF INVENTION: A POLYNUCLEIC ACID ISOLATED FROM A
TITLE OF INVENTION: PROFILE NEODODE BY THE POLYNUCLEIC ACID, A VACCINE
TITLE OF INVENTION: PREPARED FROM OR CONTAINING THE POLYNUCLEIC ACID OR
TITLE OF INVENTION: PROFIEIN,
NUMBER OF SEQUENCES: 77
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                       PEATURE: OTHER INFORMATION: Synthetic oligonucleotide with potential inhibition activity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 2.9%; Score 13; DB 1; Length 13; Best Local Similarity 100.0%; Pred. No. 1.7e+02; Matches 13; Conservative 0; Mismatches 0; Indels
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/301,435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION: 424
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/131,625
FILING DATE: 05-OCT-1993
ATTORNEY/AGENT INFORMATION:
NAME: Lavalleye, Jean-Paul M.P.
REGISTATION NUMBER: 31,451
REFERENCE/DOCKET NUMBER: 4625-021-55X CIP
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 34, Application US/08301435; Patent No. 6592873; GENERAL INFORMATION:
    Gryaznov, Sergei
Pongracz, Krisztina
                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   46 CTAACCCTAACTG 58
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Arlington
STATE: Virginia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STATE: Virginia COUNTRY: U.S.A.
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US-08-301-435-34
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APPLICANT: Matray, Tracey
ATTLE OF INVENTION: Oligonucleotide N3'-P5' Thiophosphoramidates: Their Synthesis and
FILE REFERENCE: 039/003
CURRENT APPLICATION NUMBER: U5/09/657,445A
CURRENT APPLICATION NUMBER: US 60/153,201
PRIOR APPLICATION NUMBER: US 60/153,201
PRIOR PILING DATE: 1999-09-10
PRIOR FILING DATE: 1999-10-19
NUMBER OF SEQ ID NOS: 9
SOFTWARE: Patentin Version 3.1
SEQ ID NO 8
                                                                                                                                                                                                                                                                                                                                                                            ; OTHER INFORMATION: Synthetic oligonucleotide with potential inhibition activity US-09-657-445A-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; OTHER INFORMATION: Synthetic oligonucleotide with potential inhibition activity
US-10-463-076-2
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; Sequence 8, Application US/10463076
; Patent No. 6835826
; GENERAL INFORMATION:
; APPLICANT: Geron Corporation
                                                                                                                                                                                                                                                                                           LENGTH: 13
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 2.9
Best Local Similarity 100.
Matches 13; Conservative
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                                                                                                                                                                                                                                                                                                                                                          FEATURE:
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                 2.9%; Score 13; DB 1; Length 16; 100.0%; Pred. No. 2.1e+02;
                                                                                     0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
Book ining
APPLICANT:
Stocket, Thomas A.
APPLICANT:
Stocket, Thomas A.
APPLICANT:
Folkerts, Octto
APPLICANT:
TITLE OF INVENTION:
COMPOSITION AND METHODS FOR
TITLE OF INVENTION:
TITLE OF INVEN
          Query Match 2.9%; Score 13; DB Best Local Similarity 100.0%; Pred. No. 2.1 Matches 13; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM COMDATIBLE
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/679,645
FILING DATE: July 12, 1996
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 60/001,135
FILING DATE: July 13, 1995
APPLICATION NUMBER: 08/300,726
FILING DATE: September 2, 1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READBALE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Zwick, Michael G.
APPLICANT: Edington, Brent E.
APPLICANT: McSwiggen, James A.
APPLICANT: Merlo, Patricia Ann Owens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 219/247
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
INPORMATION FOR SEQ ID NO: 828:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                      Sequence 828, Application US/08679645; Patent No. 6350934
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SECURIOR: 1263
CORRESPONDENCE ADDRESS:
ADDRESSE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
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                                                                                                                                               268 GGGCTTCTCCGG 280
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Best Local Similarity 62.5
Matches 10; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     linear
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                                                                                                                                                                                                                                                                                                                                           US-08-679-645-828
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APPLICANT: MENG, XIANG-JIN
APPLICANT: MENG, XIANG-JIN
APPLICANT: HALBUR, PARTICK G.
APPLICANT: HALBUR, PARTICK G.
APPLICANT: HOROZOV, IGOR
APPLICANT: LUM, MELISSA A.
TITLE OF INVENTION: A POLYMUCIAIC ACID ISOLATED FROM A
TITLE OF INVENTION: A PROTEIN ENCONDE BY THE POLYMUCIAIC ACID, A VACCINE
TITLE OF INVENTION: PREPARED FROM OR CONTAINING THE POLYMUCIAIC ACID OR
NUMBER OF SEQUENCES: 77
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                    Query Match 2.9%; Score 13; DB 1; Length 16; Best Local Similarity 100.0%; Pred. No. 2.1e+02; Matches 13; Conservative 0; Mismatches 0; Indels
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/10904
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION UNMBER: US 08/131,625
APPLICATION UNMBER: US 08/131,625
ATTORNEY/AGENT INFORMATION:
NAME: Lavalleye, Jean-Paul M.P.
REGISTRATION NUMBER: 31,451
REFERENCE/DOCKET NUMBER: 4625-021-55X CIP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 34, Application PC/TUS9510904
GENERAL INFORMATION:
                    TELEFAX: (703) 413-220
TELEX: 248855 OPAT UR
INFORMATION FOR SEQ ID NO: 34:
SEQUENCE CHARACTERISTICS:
LENGTH: 16 base pairs
TYPE: nucleic acid
STRANDEDNESS: unknown
                                                                                                                                                                                                                                                      TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
(703) 413-3000
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TELEFAX: (703) 413-2220
TELEEX: 248655 OPAT UR
INFORMATION FOR SEQ ID NO: 34:
SEQUENCE CHARACTERISTICS:
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PCT-US95-10904-34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         268 GGGCTTCTCCGG 280
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4 GGGCTTCTCCGG 16
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TELEPHONE:
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PCT-US95-10904-34
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US-08-770-565-2
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  Sequence 1, Application US/09220510B

Sequence 1. Application US/09220510B

GENERAL INFORMATION:
APPLICANT: RESNICK, NITZAN

TITLE OF INVENTION: EXPRESSION VECTORS COMPRISING MULTIPLE SHEAR STRESS
TITLE OF INVENTION: RESPONSIVE ELEMENTS (SSRE) AND METHODS OF USE FOR
TITLE OF INVENTION: ANGIOGENESIS IN A SHEAR STRESS ENVIRONMENT
FILE REFERENCE: P-2771-US
CURRENT APPLICATION NUMBER: US/09/220,510B

CURRENT APPLICATION NUMBER: US/09/220,510B

CURRENT PILING DATE: 1998-12-24

NUMBER OF SEQ ID NOS: 6

SOSTWARE: PALENTIN Ver. 2.1
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; Sequence 2, Application US/08770565
; Patent No. 5846723
; GENERAL INFORMATION:
    APPLICANT: Kim, Nam Woo
    APPLICANT: Kim, Nam Woo
    APPLICANT: Pruzan, Ronald
    APPLICANTION: Meintich, Scott L.
    TITLE OF INVENTION: Telomerase
    NUMBER OF SEQUENCES:
    ADDRESSEE: TOWNSEND and TOWNSEND and CREW LLP
    STREET: TWO Embarcadero Center, 8th Floor
    CITY: San Francisco
    STATE: California
    CONWTRY: USA
    COMPUTER: IBM PC compatible
    OPERATING SYSTEM: PC-DOS/MS-DOS
    SOFTWARE: PatentIn Release #1.0, Version #1.30
    CURRENT APPLICATION NORTH:
    APPLICATION NUMBER: USA
    CURSIFICATION NUMBER: 20-DEC-1996
    FILING DATE: 20-DEC-1996
    CLASSIFICATION NUMBER: ASS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ch 2.8%; Score 12.8; DB 1; Length 17; 1. Similarity 87.5%; Pred. No. 2.2e+02; 14; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: Description of Artificial sequence: OTHER INFORMATION: A PDGF-A Shear Stress Response Element.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-002300US
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     333 GGGGCGAGGCGAGG 348
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA ORGANISM: Artificial sequence
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-09-220-510B-1
US-09-220-510B-1
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US-08-770-565-2
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LENGTH: 17
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RESULT 220
3.-08-873-709-20
5. Sequence 20, Application US/08873709
5. Patent No. 6037126
6. GENERAL INFORMATION:
7. APPLICANT: Grossman, Abraham
7. TITLE OF INVENTION: APPARATUS FOR DETERMINING THE PRESENCE OF TITLE OF INVENTION: PROTEIN COMPONENT OF TELOMERASE ENZYME
7. TITLE OF SEQUENCES: 25
7. NUMBER OF SEQUENCES: 25
7. CORRESPONDENCE ADDRESS:
7. ADDRESSEE: Abraham Grossman
7. ADDRESSEE: Abraham Grossman
7. ADDRESSEE: Abraham Grossman
7. ....hington Avenue
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                                                                                                                                                     Length 30;
                                                                                                                                                                                                   Indels
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APPLICATION NUMBER: US/08/873,709 FILING DATE: 12-JUN-1997 CLASSIFICATION: 435
                                                                                                                                                     Query Match 2.8%; Score 12.8; DB 1; Best Local Similarity 70.8%; Pred. No. 2.9e+02; Matches 17; Conservative 0; Mismatches 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 12.8; DB 1;
Pred. No. 1.7e+02;
0; Mismatches 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION: 435
ATTORNEY AGENT INFORMATION:
NAME: Janiuk, Anthony J.
REGISTRATION NUMBER: 29,809
REFERENCE/DOCKET NUMBER: Q001/002
TELECHOMUNICATION INFORMATION:
TELECHOMORE: 914.747-9108
INFORMATION FOR SEQ ID NO: 20:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-09-255-464B-16; Sequence 16, Application US/09255464B; Patent No. 6238867; GENERL INFORMATION: APPLICANT: ROLLINGOR, APPLICANT: Grossman, Abraham
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET: 666 Washington Avenue
CITY: Pleasantville
STATE: NY
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 2.8%;
Best Local Similarity 70.8%;
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 62 base pairs
LENGTH: 30 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: single
                                                                   linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: 11
MOLECULE TYPE:
                                                                 TOPOLOGY: 1i
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RESULT 224
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TITLE OF INVENTION: Compositions, Methods, Kits and Apparatus for TITLE OF INVENTION: Identifying Naturally Occurring RNA Sequences Having TITLE OF INVENTION: Affinity for RNA-Binding Proteins FILE REPERENCE: Q001/004a CURRENT APPLICATION NUMBER: US/09/255,464B CURRENT FILING DATE: 1999-02-22 PRIOR APPLICATION NUMBER: 60/075,495 PRIOR FILING DATE: 1998-02-23 NUMBER OF SEQ ID NOS: 25 SOFTWARE: Patentin Ver. 2.0 SEQ ID NO 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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ZIP: 10570
COMPUTER 10570
COMPUTER: Floppy disk
COMPUTER: IBM PC COMPAID
COMPUTER: IBM PC COMPAID
COMPUTER: IBM PC COMPAID
COMPUTER: PATENTIN Release #1.0, Version #1.30
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRINT APPLICATION DATA:
APPLICATION NUMBER: US/08/873,709
FILING DATE: 12-JUN-1997
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Janiuk, Anthony J.
REGISTRATION NUMBER: 29,809
REFERENCE/DOCKET NUMBER: 29,809
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
2.8%; Score 12.8; DB 1;
Best Local Similarity 70.8%; Pred. No. 1.7e+02;
Matches 17; Conservative 0; Mismatches 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     222 TCGCCTGCCCAGCCCCGAACCCC 245
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Fatent No. 6037126
Fatent No. 6037126
Fatent INFORMATION:
APPLICANT: Grossman, Abraham
TITLE OF INVENTION: COMPOSITIONS, ITLE OF INVENTION: APPRARATUS FOR ITLE OF INVENTION: APPRARATUS FOR ITLE OF INVENTION: PROTEIN COMPONION: NUMBER OF SEQUENCES: 25
NUMBER OF SEQUENCES: 25
NUMBER OF SEQUENCES: 25
NUMBER OF SEQUENCES: 25
NUMBER OF SEQUENCES: 35
NUMBER
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I: 666 Washington Avenue
Pleasantville
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 222
US-08-873-709-19/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-09-255-464B-16
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                                                                     APPLICANT: Roninson, Igor
APPLICANT: Roninson, Igor
APPLICANT: Grossman, Abraham
TITLE OF INVENTION: Compositions, Methods, Kits and Apparatus for
TITLE OF INVENTION: Gentlifying Naturally Occurring RNA Sequences Having
TITLE OF INVENTION: Affinity for RNA-Binding Proteins
TITLE OF INVENTION: Affinity for RNA-Binding Proteins
CURRENT APPLICATION NUMBER: US/09/255,464B
CURRENT FILING DATE: 1999-02-22
RIGHT APPLICATION NUMBER: 60/075,495
PRIOR FILING DATE: 1998-02-23
NUMBER OF SEQ ID NOS: 25
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 15
LENGTH: 66
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FILING DATE: September 23, 1994
CLASSIFICATION: 435
FRIOR APPLICATION DATA:
PRIOR APPLICATION DATA: including application
PRIOR APPLICATION DATA: described
APPLICATION NUMBER: 08/008,895
FILING DATE: January 19, 1993
APPLICATION NUMBER: 07/989,849
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 658, Application US/08311486C

Sequence 658, Application US/08311486C

Pattent No. 5811300:

APPLICANT: Sean Sullivan

APPLICANT: Kevin Kisich

APPLICANT: Kevin Kisich

APPLICANT: Dan T. Stinchcomb

APPLICANT: Dan T. Stinchcomb

APPLICANT: Danes McSwiggen

TITLE OF INVENTION: RIBOZYME TREATMENT OF

TITLE OF INVENTION: RIBOZYME TREATMENT OF

TITLE OF INVENTION: RIBOZYME TREATMENT OF

TITLE OF INVENTION: TAF.'

TITLE OF INVENTION: TAF.'

CORRESPONDENCE ADDRESS:

ADDRESSEE: LYON & LYON
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COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          222 TCGCCTGCCCAGCCCCCGAACCCC 245
Sequence 15, Application US/09255464B Patent No. 6238867 GENERAL INFORMATION:
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STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 2.8°
Best Local Similarity 70.8°
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
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GENERAL INPORMATION:
APPLICANT: Shay, Jerry W.
APPLICANT: Wright, Woodring E.
APPLICANT: Platyszek, Mieczysław A.
APPLICANT: Platyszek, Mieczysław A.
APPLICANT: Platyszek, Mieczysław A.
APPLICANT: Orey, David
APPLICANT: Orey, David
APPLICANT: Orey, David
APPLICANT: Wo. 6015710cn, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by
TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 46
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisc
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 12; DB 1; Length 12;
Pred. No. 1.8e+02;
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                                             SUMTE: COUNTRY: USA

ZIP: 94111-3834

COUNTRY: USA

ZIP: 94111-3834

COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC Compatible
COMPUTER: Patentin Release #1.0, Version #1.30

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/770,565

FILING DATE: 20-DEC-1996
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944

REFERENCE/DOCKET NUMBER: 015389-002300US

TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-576-0300

INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:

'COUNTRY ALES TO THE TABLE TO 
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REFERENCE/DOCKET NUMBER: 015389-001600US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 676.0.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 2.7%; Score 12; DB Best Local Similarity 100.0%; Pred. No. 1.8 Matches 12; Conservative 0; Mismatches
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; Sequence 10, Application US/08630019A
; Patent No. 6015710
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NAME: Storella, John R.
REGISTRATION NUMBER: 32,9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          369 GGAAGAGGAACG 380
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STRANDEDNESS: single
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         San Francisco
California
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APPLICANT: Pruzan, Ronald
APPLICANT: Weinrich, Scott L.
TITLE OF INVENTION: Methods for Detecting the RNA Component of
TITLE OF INVENTION: Telomerase
NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NS-09-1228-174A-38/C

Sequence 38, Application US/09328174A

FREEDE NO. 6448003

GENERAL INFORMATION:
APPLICANT: Guida, Marco
APPLICANT: Kurth, Janice
TITLE OF INVENTION: Genotyping Human Phenol Sulfotransferase
TITLE OP INVENTION: GENOTYPING (STP2)
FILE REFERENCE: 4389-6 (formerly SEQ-16P)
CURRENT FILING DATE: 1999-06-08
PRIOR APPLICATION NUMBER: 09/328,174A
PRIOR RILING DATE: 1999-06-08
PRIOR RILING DATE: 1999-06-08
NUMBER OF SEQ ID NOS: 110
SOPTWARE: PABLESEQ for Windows Version 3.0
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Pred. No. 2.1e+02;
0; Mismatches 1; Indels
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FILING DATE: December 7, 1992
ATTORNEY/AGENT INFORMATION:
NAME: WARDING, MICHARD J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 209/166
TELECOMMUNICATION INFORMATION:
TELEPAN: (213) 489-1600
TELEFAX: (213) 955-0440
TELEFIX: 67-3510
INPORMATION FOR SEQ ID NO: 658: SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
US-08-311-486C-658
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Patent No. 5846723
GENERAL INFORMATION:
APPLICANT: Kim, Nam Woo
APPLICANT: Wu, Fred
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      302 AGAGTTGGCCTCTG 315
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Best Local Similarity 92.31
Best Local 31 Conservative
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ORGANISM: H. sapiens
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STREET: TW
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US-08-770-565-4/c
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APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
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Gaps

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Patent No. 6294650
GENERAL INFORMATION:
APPLICANT: Shay, Jerry W.
APPLICANT: Shay, Woodring E.
APPLICANT: Ocrey, David R.
APPLICANT: No. 6294650con, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
ADDRESSER: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STREET: California
                                                                            Length 12;
                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2.7%; Score 12; DB 1; Length 12; 100.0%; Pred. No. 1.8e+02; ive 0; Mismatches 0; Indels
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COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM FC Compatible

OPERATING SYSTEM: PC-DOS/NNS-DOS

SOFTWARE: DatentIn Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/349,532

FILING DATE:

CLASS!FICATION DATA:

APPLICATION NUMBER: US 08/838,545

FILING DATE: 09-APR-1996

ATTORNEY/AGENT INFORMATION:

RESTERANCY ON PROPATION:

REFERENCE SCOREIT NUMBER: 32,944

REFERENCE DOCKET NUMBER: 32,944

REFERENCE DOCKET NUMBER: 32,944

REFERENCE DOCKET NUMBER: 32,944

REFERENCE DOCKET NUMBER: 015389-001610US

TELEFONE: (415) 576-0300

INFORMATION FOR SEQ ID NO: 8

SEQUENCE CHARACTERISTICS:

LENGTH: 12 PARCE SETICS:
                                                                2.7%; Score 12; DB 1; Lk
100.0%; Pred. No. 1.8e+02;
iive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          other nucleic acid
                                                                                                              12; Conservative
                                                                                                                                                           46 CTAACCCTAACT 57
                                                                                                                                                                                      CTAACCCTAACT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 12 base pairs
TYPE: nucleic acid
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                                                              Query Match
Best Local Similarity
Matches 12; Conserv
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Best Local Similarity
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DESCRIPTION:
DESCRIPTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRANDEDNESS:
TOPOLOGY: 11
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) DESCRIPTION:
US-09-349-532-8
                                                                                                                                                                                                                                                                                          US-09-349-532-8/c
                     US-08-838-545-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY:
                                                                                                                                                                                                      12
                                                                                                                                                                                                                                                                          RESULT 229
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                                                                                                                                                              other nucleic acid (PMA), desc = "peptide nucleic acid (PMA), where (deoxy)ribose-phosphate linkages are replaced by Where (deoxy)ribose-phosphate linkages are replaced by N-(2-aminoethyl) alycine units linked to nucleotide bases via glycine amino nitrogen through a methylenecarbonyl linker"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 other nucleic acid (PNA), described acid (PNA), where (deoxy(ribose-phosphate linkages are replaced by N-(2-aminoethyl)alycine units linked to nucleotide bases via glycine amino N through a methylenecarbonyl linker.
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APPLICANT: Shay, Jerry W.
APPLICANT: Platyszek, Mieczyslaw A.
APPLICANT: Platyszek, Mieczyslaw A.
APPLICANT: Corey, David R.
APPLICANT: Ocrey, David R.
APPLICANT: No. 6046307ton, James C.
TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 60
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                       Query Match 2.7%; Score 12; DB 1; Length 12; Best Local Similarity 100.0%; Pred. No. 1.8e+02; Matches 12; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: ISM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/838,545
FILING DATE: 09-APR-1997
CLASSIPICATION: DATA:
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INPOMMATION:
NAMR: SCENEIL ACHINA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-001610US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 8, Application US/08838545
Patent No. 6046307
GENERAL INFORMATION:
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 10.
SEQUENCE CHARACTERISTICS:
LENGTH: 12 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: other nucleic
DESCRIPTION: /desc = "pept
DESCRIPTION: where (deoxy)
DESCRIPTION: where (deoxy)
DESCRIPTION: 01/ycine aminoc
                                                                                                                                                                                                                                                                                                                                                                                                                     46 CTAACCCTAACT 57
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 12 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           California
: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                linear
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DESCRIPTION:
DESCRIPTION:
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DESCRIPTION:
                                                                                                                                                                                                                                                                              US-08-630-019A-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 228
US-08-838-545-8/c
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/desc = "peptide nucleic acid (PNA),
where (deoxy(ribose-phosphate linkages are replaced by
N-(2-aminoethyl)glycine units linked to nucleotide bases via
glycine amino N through a methylenecarbonyl linker"
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                                                                                                                                                                                                                                                    STREET: Suite 4700
CITY: Los Angeles
STATE: Guite 4700
CITY: Los Angeles
STATE: 10.5 A.

ZIP: 90071-2066
COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: Storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM Compatible
OPERATING SYSTEM: IBM Compatible
OPERATING SYSTEM: IBM COMPATA:
APPLICATION NUMBER: US/08/292,620A
FILING DATE: August 17, 1994
CLASSIFICATION DATA: including application
PRIOR APPLICATION DATA: including application
PRIOR APPLICATION NUMBER: 08/008,895
FILING DATE: January 19, 1993
APPLICATION NUMBER: 07/989,849
FILING DATE: December 7, 1992
ATTORNEY AGENT INFORMATION:
REGISTRATION NUMBER: 208/149
FILING DATE: January 19, 1993
ATTORNEY AGENT INFORMATION:
REGISTRATION NUMBER: 208/149
FILING DATE: January 19, 1992
ATTORNEY AGENT INFORMATION:
REGISTRATION NUMBER: 208/149
FILING STATION NUMBER: 208/140
FILING STATION NUMBER: 208/140
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 12; DB 1; Le
Pred. No. 2.3e+02;
                                       APPLICANT: Kenneth G. Draper
TITLE OF INVENTION: BIBOZYME TREATMENT OF
TITLE OF INVENTION: BIBOZYME OF CONDITIONS
TITLE OF INVENTION: RELATED TO LEVELS OF
TITLE OF INVENTION: MOLECULE-1 (I-CAM-1)
NUMBER OF INVENTION: MOLECULE-1 (I-CAM-1)
NUMBER OF SEQUENCES: 2390
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 396, Application US/09071845

Sequence 396, Application US/09071845

Patent No. 613296,
GENERAL INFORMATION:
APPLICANT: Susan Grimm
APPLICANT: James McSwiggen
APPLICANT: Sean SUllivan
APPLICANT: Kenneth G. Draper
TITLE OF INVENTION: RIBOZYME TREATMENT OF
TITLE OF INVENTION: RELATED TO LEVELS OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 2.7%; Score 12; DB Best Local Similarity 100.0%; Pred. No. 2.3 Matches 12; Conservative 0; Mismatches
James McSwiggen
Sean Sullivan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19 CTGGGAGGGTG 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12 crescadesers 1
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                       APPLICANT:
APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                            Score 12; DB 1; Le
Pred. No. 2.3e+02;
0; Mismatches 0;
                                  Sequence 396, Application US/08292620A
Sequence 396, Application US/08292620A
GENERAL INPORMATION:
APPLICANT: Susan Grimm
APPLICANT: James McSwiggen
APPLICANT: James McSwiggen
APPLICANT: Sean Sullivan
APPLICANT: Kenneth G. Draper
ITLE OF INVENTION: RIBOZYME TRATWENT OF
ITLE OF INVENTION: RIBOZYME APPLICANT
ITLE OF INVENTION: MOLECULE-1 (I-CAM-1)
NUMBER OF SEQUENCES: 2390
CORRESPONDENCE ADDRESSS:
ADDRESSE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-08-292-620A-591/C
; Sequence 591, Application US/08292620A
; Patent No. 5837542
; GAPERAL INFORMATION:
APPLICANT: Susan Grimm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 2.7%; Scc
Best Local Similarity 100.0%; P?
Matches 12; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19 CTGGGAGGGTG 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12 crescasecerc 1
                    US-08-292-620A-396/c
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Sequence 720, Application US/09081646

Facent No. 6333152

FAREAL INFORMATION:
APPLICANT: Kinler, Kenneth
APPLICANT: Cangletein, Bert
APPLICANT: Zhang, Lin
APPLICANT: Cangletein, Bert
APPLICANT: Cangletein, Cangletein, Courage (Current Applicanto)
FILE OF INVENTY FILING DATE: 1998-05-20
FARLIER PILING DATE: 1997-05-21
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 720
LENGTH: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
2.7%; Score 12; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 2.3e+02;
Matches 12; Conservative 0; Mismatches 0;
                                                                                     COMPUTER READABLE FORM:

MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
COMPUTER: IBM COMPATIBLE
CORPATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: WORD Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/071,845
FILING DATE: CLASSIFICATION DATA:
APPLICATION NUMBER: US/08/292,620
FILING DATE: ABUGUST 17, 1994
APPLICATION NUMBER: US/08/292,620
FILING DATE: DAGUST 17, 1994
APPLICATION NUMBER: US/08/99,849
FILING DATE: December 7, 1992
APPLICATION NUMBER: 08/089,895
FILING DATE: December 7, 1993
APPLICATION NUMBER: 08/089,895
FILING DATE: December 7, 1992
APPLICATION NUMBER: 08/089,895
FILING DATE: December 7, 1992
APPLICATION NUMBER: 32,327
ATTORNEY/AGENT INFORMATION:
TELECOMMUNICATION NUMBER: 32,327
TELECOMMUNICATION NUMBER: 32,327
TELECOMMUNICATION NUMBER: 32,327
TELEFAX: (213) 955-0440
TELEX: 67,3510
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
TYPE: nucleic acid
TYPE: TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19 CTGGGAGGGTG 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12 credendedere 1
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                         COUNTRY: U.S.A.
TP: 90071-2066
                                         California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 234
US-09-081-646-720/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-09-071-845-591
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 233
US-09-071-845-591/C
Sequence 591, Application US/09071845
Fatent No. 612367
GENERAL INFORMATION:
APPLICANT: Susan Grimm
APPLICANT: James McSwigen
APPLICANT: James McSwigen
APPLICANT: Asen Sullivan
APPLICANT: Renneth G. Draper
TITLE OF INVENTION: RIBOZYME TREATMENT OF
TITLE OF INVENTION: DISEASES OR CONDITIONS
TITLE OF INVENTION: DISEASES OF CONTITIONS
TITLE OF INVENTION: INTRACELLULAR ADHESION
TITLE OF INVENTION: INTRACELLULAR ADHESION
TITLE OF INVENTION: MOLECULE-1 (I-CAM-1)
NUMBER OF SEQUENCES: 2390
CORRESONENCE ADDRESS:
TITLE OF INVENTION: INTRACELLULAR ADHESION TITLE OF INVENTION: MOLECULE-1 (I-CAM-1) NUMBER OF SEQUENCES: 2390 CORRESPONDENCE ADDRESS: ADDRESSE: Lyon & Lyon STREET: 633 West Fifth Street STREET: Suite 4700
                                                                                                                                                                      CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
                                                                                                                                                                                                                                                                                                                                         MEDIUM 11FE: SUCISAGE
COMPUTER: BEN SUCISAGE
SOFTWARE: WORD SELFECT 5.1
SOFTWARE: WORD PERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/071,845
FILING DATE: US/09/071,845
FILING DATE: Adjust 17,1994
APPLICATION NUMBER: US/08/292,620
FILING DATE: Adjust 17,1994
APPLICATION NUMBER: 0S/088,989
FILING DATE: December 7,1993
APPLICATION NUMBER: 07/989,849
FILING DATE: December 7,1992
ATTORNEY/AGENT INFORMATION:
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 32,327
TELECOMMUNICATION INFORMATION:
THE COMMUNICATION INFORMATION:
THE COMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          E: Lyon & Lyon
633 West Fifth Street
Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 396:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 100.
"...hes 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19 CTGGGAGGGGTG 30
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US-09-071-845-396
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPOSITION AND METHODS FOR MODULATION OF GENE EXPRESSION IN PLANTS
                                                                                                                                                                                                                                COUNTRY: U.S.A.

ZIP: 90071-2066

COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: etorage
COMPUTER: IDM COMPALIABLE
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: WORD PERFECT S.I.
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/679,645
FILING DATE: July 12, 1996
CLASSIFICATION: 800
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/001,135
FILING DATE: July 13, 1995
APPLICATION NUMBER: 60/001,135
FILING DATE: September 2, 1994
ATTORNEY/AGENT INFORMATION:
NAME: WAEDLUGY RICHARD J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 32,327
TELECOMMUTCATION INFORMATION:
TELECOMMUTCATION INFORMATION:
TELECOMMUTCATION INFORMATION:
APPLICANT: Folkerts, Otto
APPLICANT: Merlo, Donald J.
TITLE OF INVENTION: COMPOSITION AN
TITLE OF INVENTION: MODULATION OF
TITLE OF INVENTION: IN PLANTS
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 515:
SEQUENCE CHARACTERISTICS:
LENGTH: 16 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
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Best Local Similarity 91.7
Matches 11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                               CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: P.O. 1
CITY: Houston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ;
TOPOLOGY:
US-08-679-645-515
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                                                                                                                                                     Sequence 112, Application US/08050073

Sequence 112, Application US/08050073

GENERAL INFORMATION:
APPLICANT: Begovich, Ann B.
APPLICANT: Begovich, Ann B.
APPLICANT: Britch, Henry A.
APPLICANT: Griffith, Robert L.
APPLICANT: Griffith, Robert L.
APPLICANT: Griffith, Robert L.
APPLICANT: Stephen J.
TITLE OF INVENTION: Typing
NUMBER OF SEQUENCES: 315
CORRESPONDENCE ADDRESS:
ADDRESSEE: Hoffmann-La Roche Inc.
STREET: 340 Kingsland Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: Nutley
STATE: New Jersey
COUNTRY: U.S.A.
ZIP: 0.110
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: BAIGHTIN Release #1.0, Version #1.25
SOFTWARE: Patentin Release #1.0, Version #1.25
APPLICATION NUMBER: US/08/050,073
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Zwick, Michael G.
Edington, Brent E.
McSwiggen, James A.
Merlo, Patricia Ann Owens
Guo, Lining
Skokut, Thomas A.
Young, Scott A.
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; Patent No. 6350934
; GENERAL INFORMATION:
; APPLICANT: Zwick, Michael G.
APPLICANT: Edington, Brent E.
; APPLICANT: Merlo, Patricia Ann Owens
; APPLICANT: Guo, Lining
; APPLICANT: Skokut, Thomas A.
; APPLICANT: Skokut, Thomas A.
; APPLICANT: Young, Scott A.
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CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Petry, Douglas A.
REGISTRATION NUMBER: 35,321
REPERRICE/POCKET NUMBER: 9869
TELECOMMUNICATION:
TELEPHONE: (510) 814-2974
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: (510) 814-2974
TELEPAX: (510) 814-2977
INFORMATION FOR SEQ ID NO: 112:
SEQUENCE CHARACTERISTICS:
LENGTH: 16 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          , MOLECULE TYPE: genomic DNA
US-08-050-073-112
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Best Local Similarity 100.
                        439 GGCTCACACATG 450
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                                           12 GGCTCACATG 1
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STRANDEDNESS: singl
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                                                                                                                                         US-08-050-073-112/c
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US-08-679-645-515
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RESULT SESSUE OF SECULOR SECULOR SEQUENCE 1, Application US/08484551

Sequence 1, Application US/08484551

Sequence 1, Application US/08484551

Setent No. 5714328

APPLICANT: Sessler, Jonathan L.

TITLE OF INVENTION: RNA PHOTOCLEAVAGE USING TEXAPHYRINS

NUMBER OF SEQUENCES: 8

CORRESPONDENCE ADDRESS:

ADDRESSEE: Arnold, White & Durkee

STREET: P.O. Box 4433

STREET: P.O. Box 4433

COUNTRY: United States of America

STATE: Texas

COUNTRY: United States of America

STATE: Texas

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: IBM PC compatible

COMPUTER: Benefit Release #1.0, Version #1.30

SOFTWARE: Patentin Release #1.0, Version #1.30

SOFTWARE: Patentin Release #1.0, Version #1.30

STATE: TELING SYSTEM: US/08/484.551

FILING DATE: COCCURTENTIN Herewith

CLASSIPICATION: STATEM:

STATEMENT APPLICATION NUMBER: US/08/484.551

FILING DATE: COCCURTENTIN Herewith
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 2.6%; Score 11.8; DB 1; Length 15; Best Local Similarity 86.7%; Pred. No. 2.3e+02; Matches 13; Conservative 0; Mismatches 2; Indels
                                                              COMPUTER United States of America
ZIP: 94086
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/469,177
FILING DATE: 06-JUN-1995
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: Larson, Jacqueline S.
REFERENCE/DOCKET NUMBER: PHAY:057
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION OF 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
TYPE: nucleic acid
TYPE: ACMUNICATION LINEAR
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NAME: Parker, David L.
REGISTRATION NUMBER: 32,165
REFERENCE/DOCKET NUMBER: PHAY:047/PAR
TELECOMMUNICATION INFORMATION:
                          California
: United States of America
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "RNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         436 CTCGGCTCACACATG 450
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE: (512) 418-3000
TELEFAX: (512) 747-7577
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INFORMATION FOR SEQ ID NO: 1:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY:
                                                        COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-08-469-177-4
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APPLICANT: SESSLER, Jonathan L.
APPLICANT: IVERSON, Brent L.
APPLICANT: SANSOM, Petra I.
APPLICANT: MRIGHT, Meredith
TITLE OF INVENTION: DNA PHOTOCLEAVAGE USING TEXAPHYRINS
NUMBER OF SEQUENCES: 10
CORRESPONDENCE ADDRESS:
APDRESSEE: Pharmacvolit--
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
2.6%; Score 11.8; DB 1; Length 15;
Best Local Similarity 86.7%; Pred. No. 2.3e+02;
Matches 13; Conservative 0; Mismatches 2; Indels
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOCTWARR: Bateniln Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/310,501
FILING DATE: Concurrently herewith
CLASSIFICATION NUMBER: US 08/112,872
APPLICATION NUMBER: US 08/112,872
APPLICATION NUMBER: US 07/822,964
FILING DATE: 21-JAN-1992
APPLICATION NUMBER: US 08/227,370
FILING DATE: 11-JAN-1993
APPLICATION NUMBER: US 08/075,123
FILING DATE: 09-JUN-1994
APPLICATION NUMBER: US 07/822,964
FILING DATE: 01-JAN-1992
APPLICATION NUMBER: US 07/771,393
FILING DATE: 01-JAN-1992
APPLICATION NUMBER: US 07/771,393
FILING DATE: 01-JAN-1992
APPLICATION NUMBER: US 07/771,393
FILING DATE: 11-JAN-1990
APPLICATION NUMBER: US 07/771,393
FILING DATE: 01-JAN-1990
APPLICATION NUMBER: US 07/320,293
FILING DATE: 01-JAN-1990
APPLICATION NUMBER: US 07/320,293
FILING DATE: 01-JAN-1990
APPLICATION NUMBER: US 07/320,293
FILING DATE: 01-JAN-1990
ATTORNEY/AGENT INFORMATION:
NAME: PATEREY DAY: 034/PAR
TELECOMMUNICATION NUMBER: PHAY: 034/PAR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               E: Pharmacyclics, Inc.
995 East Argues Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 238
US-08-469-177-4/c
; Sequence 4, Application US/08469177
; Patent No. 5607924
; PATENT INFORMATION:
APPLICANT: MAGDA, Darren
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear _
MOLECULE TYPE: RNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                436 CTCGGCTCACACATG 450
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INFORMATION FOR SEQ ID NO: 6
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE: 512/418-3000
TELEFAX: 512/474-7577
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15 ccceecrcacaeare 1
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MOLECULE TYPE:
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COMPUTER: IB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-08-323-192D-11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ò
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Sequence 5, Application US/08484551

GENERAL INFORMATION:
APPLICANT: Magda, Darren
APPLICANT: Seaeler, Jonathan L.
TITLE OF INVENTION: RNA PHOTOCLEAVAGE USING TEXAPHYRINS
NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
ADDRESSEE: Arnold, White & Durkee
STREET: P.O. Box 4433
CITY: Houston
STATE: Texas
COUNTRY: United States of America
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2.6%; Score 11.8; DB 1; Length 15;
86.7%; Pred. No. 2.3e+02;
tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                              Query Match 2.6%; Score 11.8; DB 1; Length 15; Best Local Similarity 86.7%; Pred. No. 2.3e+02; Matches 13; Conservative 0; Mismatches 2; Indels
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/484,551
FILING DATE: Concurrently herewith CLASSITCATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: Parker, David L.
REGISTRATION NUMBER: PHAY:047/PAR
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION INFOR
; SEQUENCE CHARACTERISTICS:
; LENGTH: 15 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: other nucleic acid
; DESCRIPTION: /desc = "DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: other nucleic acid;
BESCRIPTION: /desc = "RNA"
US-08-484-551-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC_DOS/MS_DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; Sequence 18, Application US/08486962
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      436 CTCGGCTCACATG 450
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Best Local Similarity 86.7
Matches 13, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 linear
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US-08-486-962-18/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2.6%; Score 11.8; DB 1; Length 15; 86.7%; Pred. No. 2.3e+02; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Peter
RECOMBINANT NEGATIVE STRAND RNA VIRUS
EXPRESSION SYSTEMS AND VACCINES
                                                                                                                                                                                                                                                                                                                                                   METHOD OF PHOSPHATE ESTER HYDROLYSIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE PATENTIN ROLES #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/486,962 FILING DATE: US/08/1895 FILI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 11, Application US/08323192D
Patent No. 5786199
GENERAL INFORMATION:
TITLE OF INVENTION: RECOMBINANT NEGATIVE
TITLE OF INVENTION: REPRESSION SYSTEMS AN
TUMBER OF SEQUENCES: 70
CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
CITY: New York
STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM : WEDIUM TYPE: Floppy disk COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESBEE: Pharmacyclics, Inc. STREET: 995 E. Arques Avenue CITY: Sunnyvale STATE: California
                                Sessler, Jonathan L.
Wright, Meredith
Ross, Kevin L.
                                                                                                              APPLICANT: ROSS, Kevin L.
APPLICANT: Miler, Richard A.
APPLICANT: Discourse A.
APPLICANT: Smith, Uadimir A.
APPLICANT: Smith, Daniel A.
TITLE OF INVENTION: METHOD OF P.
UNDRER OF SEQUENCES: 18
CORRESPONDENCE ADDRESS:
ADDRESSEE: Pharmacyclics, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              E: Floppy disk
IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           436 CTCGGCTCACACATG 450
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ccceecrcacagare 1
Magda, Darren
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Best Local Similarity 86.7
Matches 13, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 15 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 10036-2711
COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: USA
ZIP: 94086-4521
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Gaps
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                                                                                                                                                                       Query Match 2.6%; Score 11.8; DB 1; Length 15; Best Local Similarity 86.7%; Pred. No. 2.3e+02; Matches 13; Conservative 0; Mismatches 2; Indels
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2.6%; Score 11.8; DB 1; Length 15;
Best Local Similarity 66.7%; Pred. No. 2.3e+02;
Matches 10; Conservative 3; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITAIN: SULLEY 4700
CITY: LOS Angeles
STATE: California
COUNTY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
COMPUTER: BM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: WORD PERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/311,486C
FILING DATE: September 23, 1994
CLASSIFICATION DATA:
PRIOR APPLICATION NUMBER: 08/008,895
FILING DATE: December 7, 1992
APPLICATION NUMBER: 07/989,849
FILING DATE: December 7, 1992
ATTORNEY AGENT INPERMATION:
                                                                                                                                                                                                                                                                                                                                                                               Sequence 218, Application US/08311486C
; Sequence 218, Application US/08311486C
; Patent No. 5811300
; GENERAL INFORMATION:
APPLICANT: Sean Sullivan
APPLICANT: Kevin Kisich
APPLICANT: Revin Kisich
APPLICANT: Dan T. Stinchcomb
APPLICANT: James McSwiggen
TITLE OF INVENTION: RIBOZYME TREATMENT OF
TITLE OF INVENTION: RIBOZYME
TITLE OF INVENTION: RIBASES OR CONDITIONS
TITLE OF INVENTION: RIPA
TITLE OF INVENTION: TIPP.
NUMBER OF SQUENCES: 1157
ADDRESSES: Lyon & 1...-
STREEP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 209/166
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-25-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                373 GAGGAACGGAGCGAG 387
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                                      TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEX: 67.3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
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STRANDEDNESS: single
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TOPOLOGY: ]
US-08-311-486C-218
                                                                                           ; TOPOLOGY:
US-08-294-424-35
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 35, Application US/08294424
Patent No. 5800984
Patent No. 5800984
Patent No. 5800984
Patent No. 5800984

APPLICANT: Vary, Calvin
TITLE OF INVENTION: NUCLEIC ACID SEQUENCE DETECTION BY TITLE OF INVENTION: TRIPLE HELIX FORMATION
NUMBER OF SEQUENCES: 49
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson
STREET: 225 Franklin Street
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/323,192D
FILING DATE: 14-OCT-1994
ILLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: COLUZZI, LAURA A.
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 7682-035
TELECOMMUNICATION INFORMATION:
TELECHONICATION TOR SEQ 1D NO: 11:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDENSS: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SIALE: WASBRACHBELTS
COUNTYEY: U.S.A.
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage
COMPUTER: IBM PS/2 Model 502 or 55SX
OPERATING SYSTEM: IBM P.C. DOS (Version 3.30)
SOFTWARE: WordPerfect (Version 5.0)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/294,424
FILING DATE:
CLASSIFICATION NUMBER: US/08/294,424
FILING DATE: 10.704 1993
APPLICATION NUMBER: US/08/000,922
FILING DATE: 10.704 1993
APPLICATION NUMBER: US/07/629,601B
FILING DATE: 17-DEC-1990
ATTORNEY/AGENT INFORMATION:
NAME: Freeman, John W.
REGISTRATION NUMBER: 29,066
REFERENCE/DOCKET NUMBER: 29,066
REFERENCE/DOCKET NUMBER: 29,066
REFERENCE/DOCKET NUMBER: 29.066
REFERENCE/DOCKET NUMBER: 29.066
RELECOMMUNICATION INFORMATION:
TELLEPOWE: (617) 542-8906
TELLEPAX: 200154
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEX: 200154
INFORMATION FOR SEQ ID NO: 35:
SEQUENCE CHARACTERISTICS:
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STATE: Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; TOPOLOGY: unknown
; MOLECULE TYPE: RNA
US-08-323-192D-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-08-294-424-35/c
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US-08-316-439A-8

| Sequence 8, Application US/08316439A
| Sequence 8, Application US/08316439A
| Patent No. 5840520
| GENERAL INFORMATION:
| APPLICANT: CLARKE, DAVID KIRKWOOD
| TITLE OF INVENTION: SYSTEMS
| VINBER OF SEQUENCES: 43
| CORRESPONDENCE ADDRESS: ADDRESSE: COOLEY GODWARD CASTRO HUDDLESON & TATUM
| STREET: FIVE PALO ALTO
| STREET: FIVE PALO ALTO
| STATE: CALIFORNIA
| COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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                                                                                                                                          STREET: SUITCE 4/700
CITY: LOS ANGELES
COUNTRY: LOS ANGELES
ZIP: GOIL-2066
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 1BM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA: W/08/292,620A
FILING DATE: August 17, 1994
CLASSIFICATION DATA: including application
PRIOR APPLICATION DATA: described below:
PRIOR APPLICATION DATA: including application
PRIOR APPLICATION DATA: including application
PRIOR APPLICATION DATA: 1993
APPLICATION NUMBER: 08/008 895
FILING DATE: December 7, 1992
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 32,327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 34306
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
2.6%; Score 11.8; DB 1;
Best Local Similarity 86.7%; Pred. No. 2.38+02;
Matches 13; Conservative 0; Mismatches 2;
                                                                ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    292:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21 GGGAGGGGTGGTGGC 35
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
NUMBER OF SEQUENCES: 2.
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-08-292-620A-292
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                                                                                                                                                                                                   RESULT 245

US-00-470-887A-10

Sequence 10. Application US/08470887A

Petent No. 5820871

GENERAL INFORMATION:

APPLICANT: Palese, Peter

APPLICANT: Garcia-Seatre, Adolfo

TILE OF INVENTION: EXPRESSION SYSTEMS AND VACCINES

TILLE OF INVENTION: EXPRESSION SYSTEMS AND VACCINES

NUMBER OF SEQUENCES: 60

CORRESPONDENCE ADDRESS:
ADDRESSES: Pennie & Edmonds

STREET: 1155 Avenue of the Americas

CITY: New York

COUNTRY: USA

ZIP: 10036-2711

COMPUTER READBALE FORM:
MEDIUM TYPE: FLOPPY disk

COMPUTER: IBM PC compatible

COMPUTER: IBM PC compatible

COMPUTER: BEATCHIN Release #1.0, Version #1.25

COMPUTER: Patentin Release #1.0, Version #1.25

COMPUTER: DEC JOS/MS-DOS

SOFTWARE: WESTERIED SYSTEM: US/08/470,887A

FILING DATE: 06-JUN-1995

CURSSIPTCATION NUMBER: US/08/40,887A

FILING DATE: 06-JUN-1995

TELEPHONE: (212) 790-909

TELEPHONE: (212) 869-91/8864

TELERAX: (212) 869-91/8864

TELERAX: (212) 869-91/8864

TELERAX: (212) 869-91/8864

TELERAX: 115 DASS PAILS

LUNDORMATION POR SEQ ID NO: 10:

SEQUENCE CHARACTERISTICS:

TURCITH: 15 DASS PAILS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NS-08-292-620A-292/C

Sequence 292, Application US/08292620A

Patent No. 5837542

GRNERAL INFORMATION:
APPLICANT: Susan Grimm
APPLICANT: James McSwiggen
APPLICANT: James McSwiggen
APPLICANT: Sean Sullivan
APPLICANT: Reneth G. Draper
ITLE OF INVENTION: RIBOZYME TREATMENT OF
ITLE OF INVENTION: RELATED TO LEVELS OF
ITILE OF INVENTION: INTRACELLULAR ADHESION
ITILE OF INVENTION: MOLECULE-1 (1-CAM-1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: RNA (genomic)
                                      436 CTCGGCTCACACATG 450
                                                                      1 CUUGGCUCAGACAUG 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   72 CGCCGTGCTTTTGCT 86
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STRANDEDNESS: singl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
Matches 7; Conserva
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2.6%; Score 11.8; DB 1; Length 15;
46.7%; Pred. No. 2.3e+02;
tive 6; Mismatches 2; Indels
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| Patent No. 5861244
| GENERAL INFORMATION:
| APPLICANT: WANG, C. -G.
| APPLICANT: HEPBIRN, A. G.
| TITLE OF INVENTION: GENERIC SEQUENCE ASSAY USING DNA TITLE OF INVENTION: GENERIC SEQUENCE ASSAY USING DNA TITLE OF INVENTION: TRIPLE-STRAND FORMATION:
| CORRESPONDENCE ADDRESS: 365
| ADDRESSEE: PROFILE DIAGNOSTIC SCIENCES, INC., STREET: 510 EAST 73RD STREET,
| CITY: NEW YORK | STATE: NE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear
MOLECULE TYPE: genomic DNA
DESCRIPTION: hepatitis B virus adr isolate,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CALF: 10021.
COMPUTER: 3.5 inch, 1.44Mb storage
COMPUTER: 1BM FO.XT/AT
COMPUTER: 1BM FO.XT/AT
COMPUTER: 1BM FO.XT/AT
COMPUTER: 1BM FO.MS-DOS version 6.2
SOFTWARE: Wordperfect Version 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/173,489C
FILING DATE: 22 DEC 1993
CIASSIFICATION NUMBER: US/08/436
FILING DATE: 29 OCT 1992
APPLICATION NUMBER: US 07/968,436
FILING DATE: 29 OCT 1992
ATTORNEY/AGENT INFORMATION:
NAME: Handelman, Joseph H.
REGISTRATION NUMBER: 26,179
REFERENCE/DOCKET NUMBER: U9518-6
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ey) (212) 708-1880
(212) 246-8959
O: 141:
                                                                              FILING DATE: 01-UN-1994
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: COLUZZI, LAURA A.
REGISTRATION NUMBER: 30,742
REPERENBE/DOCKET NUMBER: 7662-034
TELEPHONE: (212) 790-9090
TELEPHONE: (212) 790-9090
TELEPKA: (212) 869-9741/8864
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: NUCleic acid
STRANDENESS: single
                                  JMBER: US/08/252,508B
01-JUN-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: double stranded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; TOPOLOGY: unknown
; MOLECULE TYPE: RNA (genomic)
US-08-252-508B-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     72 CGCCGTGCTTTGCT 86
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 46.74
Matches 7; Conservative
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                                                 APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 249
US-08-173-489C-141/c
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US-08-252-508B-10

i Sequence 10, Application US/08252508B

i Patent No. 5654037

i GENERAL INFORMATION:

APPLICANT: Patese, Peter
APPLICANT: Garcia-Sastre, Adolfo
ITILE OF INVENTION: EXPRESSION SYSTEMS AND VACCINES
ITILE OF SEQUENCES: 60

CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
ITILE OF UNIVENTION: ADDRESSEE:
ADDRESSEE: New York
STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2.6%; Score 11.8; DB 1; Length 15; 46.7%; Pred. No. 2.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2; Indels
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ZIP: 10036-2711
COMPUTER READABLE PORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/316,439A
FILING DATE: September 30, 1994
CLASSIFICATION 1424

RRIOR APPLICATION NUMBER: 081190,678
FILING DATE: February 1, 1994
CLASSIFICATION NUMBER: 07/925,061
FILING DATE: August 4, 1992
CLASSIFICATION NUMBER: 07/925,061
FILING DATE: May 22, 1990
CLASSIFICATION NUMBER: 07/527,237
FILING DATE: May 22, 1990
CLASSIFICATION NUMBER: 07/527,237
FILING DATE: May 22, 1990
CLASSIFICATION NUMBER: 07/440,053
FILING DATE: No. 5840520ember 21, 1989
CLASSIFICATION NUMBER: 07/399,728
FILING DATE: August 28, 1989
CLASSIFICATION: 424
PRIOR APPLICATION: 424
PRIOR APPLICATION: 424
FILING DATE: No. 5840520ember 21, 1989
CLASSIFICATION: 424
ATTORNEY, AGENT INFORMATION:
FILING DATE: August 28, 1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      6; Mismatches
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MOLECULE TYPE: SYNTHETIC DNA
US-08-316-439A-8
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Best Local Similarity 46.73
Matches 7; Conservative
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TYPE: nucleic acid
STRANDEDNESS: single
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0; Gaps

Gaps ô

Indels

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Mismatches
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CLASSIPECATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/252,508
FILING DATE: 01-010-1994
ATTORNEY/AGENT INFORMATION:
NAME: Coruzzi, Laura A.
REGISTATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 7682-034
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-9741/8864
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INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: RNA (genomic)
US-09-106-377-10
                                                    389 CCCCGCGCGCGCGC 403
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13; Conservative
                                                                                      TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-09-071-845-292/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY:
  Matches
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US-08-550-120-3
US-08-550-120-3
Sequence 3, Application US/08550120
Patent No. 598554
GENERAL INFORMATION:
APPLICANT: Hiroshi TAMIMURA et al.
APPLICANT: Hiroshi TAMIMURA et al.
TITLE OF INVENTION: METHOD FOR PROBING THE FUNCTION OF A PROTEIN
NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
ADDRESSEE: Wenderch, Lind & Ponack
STREET: 805 Fifteenth Street, N.W., #700
CITY: Washington
STREET: BOS Fifteenth Street, N.W., #700
CITY: Washington
STATE: D.C.
COUNTRY: U.S.A.
ZIP: ZOO.
STATE: D.C.
COUNTRY: U.S.A.
ZIP: MEDIUM TYPE: Diskette, 3.5 inch, 1.44 mb
COMPUTER: IBM Compatible
OPERATING SYSTEM: MS-DOS
SOFTWARE: Wordperfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/550,120
FILING DATE: October 30, 1995
CLASSIFICATION NUMBER: US/08/550,120
FILING DATE: No. 588554ember 2, 1994
ATTORNEY, AGENT INFORMATION:
REGISTRATION NUMBER: 33,367
REFERENCE/DOCKET NUMBER:
TELECOMMUNICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                           ONGANISM: Hepatitis B virus

INDIVIDUAL ISOLATE: adr

PRELICATION INFORMATION:
AUTHORS: Yoleyama, T, Ohromo, N, Matsubara, K.
AUTHORS: Yoneyama, T, Ohromo, N, Matsubara, K.
TITLE: Cloning and structural
TITLE: analysis of Hepatitis B virus DNAs subtype adr
JOHNNAL: Nucleic Acids Research
VOLUME: 11
PAGES: 4601-4610
DATE: 1983
RELEVANT RESIDUES IN SEQ ID NO: 141:FROM 1 TO 15
US-08-173-489C-141
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     2.6%; Score 11.8; DB 1;
86.7%; Pred. No. 2.3e+02;
tive 0; Mismatches 2;
  nucleotides 2405 to 2419
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              373 GAGGAACGGAGCGAG 387
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Best Local Similarity 86.79
....hes 13; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
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DESCRIPTION: nu
HYPOTHETICAL: no
ANTI-SENSE: no
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-08-550-120-3
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RESULT 251
US-09-106-377-10

$ Sequence 10, Application US/09106377

$ Patent No. 6001634

$ GENERAL INFORMATION:

$ APPLICANT: Patese, Peter

APPLICANT: Garcia-Sastre, Adolfo

TITLE OF INVENTION: RECOMBINANT NEGATIVE STRAND RNA VIRUS

$ TITLE OF INVENTION: EXPRESSION SYSTEMS AND VACCINES

NUMBER OF SEQUENCES:

$ ADDRESSEE: Pennie & Edmonds

$ STREET: 1155 Avenue of the Americas

$ STREET: New York

$ STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                            Query Match
2.6%; Score 11.8; DB 1;
Best Local Similarity 46.7%; Pred. No. 2.3e+02;
Matches 7; Conservative 6; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence, 292, Application US/09071845
Fatent No. 6132967
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Susan Grimm
APPLICANT: Dan T. Stinchcomb
APPLICANT: James McSwiggen
APPLICANT: Sean Sullivan
APPLICANT: Kenneth G. Draper
TITLE OF INVENTION: RIBOZYME TREATMENT OF
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OTHER INFORMATION: Description of Artificial Sequence:other nucleic
OTHER INFORMATION: acid
US-09-180-437-212
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
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86.7%; Pred. No. 2.3e+02;
tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 2.6%; Score 11.8; DB 1; Length 15; Best Local Similarity 86.7%; Pred. No. 2.3e+02; Matches 13; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2; Indels
     COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE 3.5 INCH 1.44 MB STORAGE
COMPUTER: IBM COMPATIBLE
COMPUTER: IBM COMPATIBLE
COMPUTER: IBM COMPATIBLE
COMPUTER: MONDERRECT 6.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/871,732A
FILING DATE:
CLASSIFICATION DATA:
ATTORNEY, AGENT INFORMATION:
NAME: MCGREGOR, MARTIN L.
REGISTRATION NUMBER: 29,329
REFERENCE/DOCKET NUMBER: 1-1
TELECHMONICATION INFORMATION:
TELEFHONE: 713-682-1213
TELEFERX: 713-682-1213
TELEFERX: 713-682-5807
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 212, Application US/09180437
Fatent No. 6251873
GENERAL INFORMATION:
APPLICANT: PUKUSAWA, Yoshichi
APPLICANT: WORISAWA, Yoshichi
TITLE OF INTENTION:
TITLE OF INTENTION: Antisense Compounds to CD14
TITLE OF INTENTION: Antisense Compounds to CD14
TITLE REPERRUCE: 1110-209P
CURRENT APPLICATION NUMBER: US/09/180, 437
CURRENT FILING DATE: 1998-11-06
EARLIER FILING DATE: 1998-03-09
EARLIER FILING DATE: 1998-03-09
EARLIER FILING DATE: 1998-03-09
SARLIER FILING DATE: 1998-03-09
SOFTWARE: PARENTIN VOS: 289
SOFTWARE: PARENTIN VOY: 2.0
SERVING DINO 212
                                                                                                                                                                                                                                                                                                                                                                                             TELEFACK: 713-682-1213
TELEFACK: 713-682-5807
TELEX: NONE
INPORMATION FOR SEQ 1D NO: 9:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 BASE PAIRS
TYPE: NUCLEIC ACID
STRANDEDNESS: SINGLE
TOPOLOGY: LINEAR
MOLECULE TYPE: OTHER NUCLEIC ACID
HYPOTHETICAL: NO
HYPOTHETICAL: NO
US-08-871-732A-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21 GGGAGGGGTGGTGGC 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  264 GCCCGGGGCTTCTCC 278
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Best Local Similarity 86.73
Matches 13, Conservative
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US-09-180-437-212/c
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Patent No. 6146074
GENERAL INFORMATION:
APPLICANT: 0'BRIEN, TIMOTHY J.
APPLICANT: WANG, YIN
TILE OF INVENTION: CELLS, VECTORS AND EXPRESSION METHOD FOR PRODUCING THE NOVEL
TITLE OF INVENTION: PROTEIN, ANTIBODIES AND USES
NUMBER OF SEQUENCES: 16
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   .;
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          DISEASES OR CONDITIONS
RELATED TO LEVELS OF
INTRACELLULAR ADHESION
MOLECULE-1 (I-CAM-1)
TITLE OF INVENTION: DISEASES OR CONDITIC
TITLE OF INVENTION: RELATED TO LEVELS OF
TITLE OF INVENTION: INTRACELLULAR ADHEST
TITLE OF INVENTION: INTRACELLULAR ADHEST
TITLE OF INVENTION: MOLECULE-1 (I-CAM-1)
NUMBER OF SEQUENCES: 2390
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STREET: Los Angeles
STREET: U.S.A.
ZIP: 90071-2066
COMPUTER FRADABLE FORM:
MEDIUM TYPE: Storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:

PAPLICATION NUMBER: US/09/071,845
FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/292,620
FILING DATE: August 17, 1994.
APPLICATION NUMBER: US/08/393
FILING DATE: December 7, 1992
FILING DATE: December 7, 1992
ATTORNEY DECEMBER: 07/989,849
FILING DATE: December 7, 1992
ATTORNEY NUMBER: 07/989,449
FILING DATE: December 7, 1992
ATTORNEY REPRENTION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 208/149
TELECOMMUNICATION INFORMATION:
TELEFRAX: (213) 489-1600
TELEFAX: 67-3510
TELEFAX: 67-3510
SEQUIBENCE CHARACTERISTICS:
TENNOTH: 15 hase pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: MARTIN L. MCGREGOR
STREET: 5380 WEST 34TH STREET, #345
CITY: HOUSTON
                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: WORD Perfect 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STATE: TEXAS
COUNTRY: UNITED STATES OF AMERICA
ZIP: 77092
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21 GGGAGGGGTGGTGGC 35
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TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              13; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear
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US-08-871-732A-9
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Query Match 2.6%; Score 11.8; DB 1; Length 15; Best Local Similarity 86.7%; Pred. No. 2.3e+02; Matches 13; Conservative 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Patent No. 5166057
, APPLICANT: PALESE, PETER; PARVIN, JEFFREY D.; KRYSTAL, MARK
, TITLE OF INVENTION: RECOMBIANT NEGATIVE STRAND RNA VIRUS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
2.6%; Score 11.8; DB 1;
Best Local Similarity 46.7%; Pred. No. 2.38+02;
Matches 7; Conservative 6; Mismatches 2;
                                                                                                          ; TOPOLOGY: linear SEQ ID NO: 106: US-09-544-934B-106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EXPRESSION-SYSTEMS

NUMBER OF SEQUENCES: 43

CURRENT APPLICATION DATA:

FILING DATE: 22-MAY-1990

PRICH APPLICATION NUMBER: 440,053

PRICH APPLICATION NUMBER: 440,053

FILING DATE: 21-MOV-1989

PILING DATE: 28-MUG-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Search completed: August 24, 2005, 14:29:01 Job time : 3 secs
INFORMATION FOR SEQ ID NO: 106:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 basepairs
TYPE: nucleic acid basepairs
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                            180 CAGCTGCTGGCCCGT 194
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Tina Anderson Hollerup
TILE OF INVENTION: No. 6753421el Process For The Detection of Mycobacteria
NUMBER OF SEQUENCES: 123
CORRESPONDENCE ADDRESS:
ADDRESSE: FINNEGAN, HENDERSON, FARABOW, GARRETT, & DUNNER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                         Sequence 9, Application US/09346510B

Sequence 9, Application US/09346510B

Batent No. 6281014

GENERAL INFORMATION:

APPLICANT: O'Brien, Timothy J.

APPLICANT: Wang, Yinxiang

TITLE OF INVENTION: SH3-Containing Protein, DNA and Uses Thereof

TITLE OF INVENTION: LOS21CIP

CURRENT APPLICATION NUMBER: US/09/346,510B

CURRENT FILING DATE: 1999-07-01.

PRIOR APPLICATION NUMBER: 08/871,732

PRIOR APPLICATION NUMBER: 08/871,732

NUMBER OF SEQ ID NOS: 32

SEQ ID NO 9

LENGTH: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ) OTHER INFORMATION: nucleotide sequence of clone 17 isolated using the ) OTHER INFORMATION: CASTING approach US-09-346-510B-9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:

MEDIUM TYPE: PLOBPY disk 3.5 inch
COMPUTER: IBM PC compatible
OPERATING SYSTEM: ASCXI
SOFTWARE: MICROSOFE Word
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/544,934B
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/029,595
FILING DATE: 15-OCt-96
APPLICATION NUMBER: 60/029,595
FILING DATE: 15-OCt-96
APPLICATION NUMBER: 60/029,595
FILING DATE: 23-OCT-96
APPLICATION NUMBER: 60/045,962
FILING DATE: 08-May-97
APPLICATION NUMBER: 08/943,777
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: Washington
STATE: District of Columbia
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Anthony C. Tridico
REGISTRATION NUMBER: 45,958
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 408-4173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 106, Application US/09544934B; Patent No. 6753421; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEPHONE: (202) 408-4173
TELEFAX: (202) 408-4400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: 1300 I ST. NW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Henrik Stender
                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21 GGGAGGGTGGTGGC 35
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US-09-544-934B-106
                            RESULT 255
US-09-346-510B-9
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GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compugen Ltd. ic search, using sw model gust 24, 2005, 14:36:10; Search time 2 Sec. 3.48 Million cell	436-060A-16 ttgcggagggtgggcct TTY_NUC 10.0 , Gapext 0.5 ags, 7423 residues	nits satisfying chosen parameters: nngth: 0 ength: 2000000000 Minimum Match 0% Maximum Match 100% Listing first 376 summaries	rnpb.subdb:* No. is the number of results predicted by chance to ha greater than or equal to the score of the result being: derived by analysis of the total score distribution. SUMMARIES Query Descrit	9 31 1 US-10-714-195-84 9 31 1 US-10-310-812-87 1 1 US-10-310-812-27 1 30 1 US-09-9057-351-22 1 30 1 US-10-359-935-22 1 30 1 US-10-359-935-22 1 30 1 US-10-310-812-4 1 30 1 US-10-811-033-4 1 30 1 US-10-811-033-5 1 0S-10-811-033-5 1 US-10-811-033-5 1 US-10-811-033-5 1 US-10-811-033-3 1 US-10-811-033-3 1 US-10-811-033-3 1 US-10-811-033-3 1 US-10-811-033-3 1 US-10-811-033-3 1 US-10-945-606-26 1 US-10-044-692-312 1 US-10-044-692-312 1 US-10-044-692-312 1 US-10-359-935-30 1 US-10-359-935-30 1 US-10-359-935-30 1 US-10-877-124-598 1 US-10-877-124-598 1 US-10-877-124-598 1 US-10-877-124-598 1 US-10-877-124-598 2 US-10-877-124-598 3 US-10-877-124-598 3 US-10-877-124-598 3 US-10-877-124-598 3 US-10-877-124-598 3 US-10-877-124-598 4 US-10-877-124-598 5 US-10-878-124 5
Copy OM nucleic - nucleic s Run on: August	US-0 t score: 451 nce: 1 gg ng table: IDEN Gapo	Total number of hits some minimum DB seq length: Maximum DB seq length: Post-processing: Minimum Dest-processing: Minimum Listir	Database : rnpb. Pred. No. is the score greater the and is derived the score greater the and is derived the score Match No. Score Match	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0

US09436060A-16.rnpb.sl

Sequence 595, App Sequence 596, App Sequence 597, App Sequence 598, App Sequence 599, App	9 9	99	9	9 6	15	9 5	619	61	9	9	62	62	62	9 0	6 2	62	62	62	63	2 6	63.5	63	63	63	9 6	6.4	quence 12	e 54	equence 54	, 4 1	ence 13	12	7	13	מ קינ	, ה ה	9,7	315	4	4	Sequence 4, Appli	14.	146	75(750	49	4 ° C	2.5	27,0	3581,	5, App	336	1471,
US-10-923-330-595 US-10-923-330-596 US-10-923-330-597 US-10-923-330-598 US-10-923-330-599	US-1	US-1	US-1 US-1	US-1 US-1	US-1	US-1	US-1	US-1	US-1	US-1	US-1	US-1	us-1	1-SU 11S-1	US-1	US-1	US-1	US-1	18-1	US-1	US-1	US-1	US-1	US-T	US-10	US-1	US-1(US-1(SO L	US-1(US-0	US-1(US-1(OT-SO	110-11	118-11	US-10	US-10	US-10	US-10	US-10	0S-05	0S-05	US-10	US-10	T-SD	118-10	US-10	US-09		1-930-423-5	-930-423-336	-930-423-
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US-09-827 US-09-746 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745 US-09-745	S-10-96 SS-10-96 SS-10-51 SS-10-51 SS-10-72 SS-10-72 SS-09-78 SS-09-78 SS-09-78 SS-09-78 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-09-84 SS-0
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Gaps
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13;
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MEDIUM TYPE: Floppy disk
COMPUTER: Eloppy disk
COMPUTER: El PR PC Compatible
COMPUTER: El PR PC Compatible
COMPUTER: El PR PC Compatible
COMPUTER: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION NAMER: US/09/057,351
FILING DATE: 08-APR.1994
CLASSIFICATION NAMER: US 08/272,102
FILING DATE: 07-JU1-1994
FILING DATE: 07-JU1-1994
PRIOR APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION NUMBER: US 08/472,802
FILING DATE: 07-JUN-1955
ATONINEY/AGBNT INFORMATION:
NAME: SECORELLA, JOHN REGION NUMBER: 32,944
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor STATE: San Francisco STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; Sequence 22, Application US/09057351
; Patent No. US20010034439A1
; GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junia
APPLICANT: Funk, Walter
TITLE OF INVENTION: Mammalian Telomerase
; TITLE OF INVENTION: Mammalian Telomerase
; UNWBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                42 TIGICTAACCCTAACTGAGAAGGGCGTAGGC 72
                         TITLE REFERENCE: 011/006C
CURRENT APPLICATION NUMBER: US/10/811,033
CURRENT FILING DATE: 2004-03-26
FURDENT FILING DATE: 2004-03-26
PRIOR PELICATION NUMBER: US/10/330,872A
PRIOR FILING DATE: 2002-12-24
PRIOR APPLICATION NUMBER: 08/510,736
PRIOR PILING DATE: 1995-08-04
PRIOR PLING DATE: 1995-08-04
PRIOR APPLICATION NUMBER: 08/420,056
PRIOR PLING DATE: 1997-01-18
PRIOR APPLICATION NUMBER: 09/420,056
PRIOR PLING DATE: 1999-10-18
PRIOR PLING DATE: 2000-11-20
NUMBER OF SEQ ID NOS: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   6.9%; Score 31; DB 100.0%; Pred. No. 13; iive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            015389-000821US
                    TITLE OF INVENTION: Inhibitors
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TELECOMMUNICATION INFORMATION: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity 100.0
Matches 31; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: USA
ZIP: 94111-3834
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                                                                                                                                                                                                                                                                                                                                                                                                       SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 1, Application US/10330872

Sequence 1, Application US/10330872

Publication No. US200301862821

GENERAL INFORMATION:

APPLICANT: Geron Corporation

APPLICANT: Meinrich, Scott

APPLICANT: Akinson III, Edward

APPLICANT: Vasserot, Alain

APPLICANT: Ussuer, Road

TITLE OF INVENTION: Unhibitors

FILE REPREBRENCE: 101/006C

CURRENT APPLICATION NUMBER: 08/10/330,872

CURRENT FILING DATE: 1995-04-04

FRIOR FILING DATE: 1995-04-04

FRIOR FILING DATE: 1995-04-04

FRIOR FILING DATE: 1995-10-18

FRIOR FILING DATE: 2000-11-20

NUMBER OF SEQ ID NOS: 11

SED IN NO. 11-20

SED TWARE: PATENTIN VERSION 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 1, Application US/10811033
| Publication No. US2005008983A1
| GENERAL INFORMATION:
| APPLICANT: Geron Corporation
| APPLICANT: Atkinson III, Edward
| APPLICANT: Lichtsteiner, Serge
| APPLICANT: Lichtsteiner, Serge
| APPLICANT: Pruzan, Ronald
| APPLICANT: Pruzan, Ronald
| TITLE OF INVENTION: Using Purified Telomerase to Identify Telomerase Activators and
                                                                                                                                                                                                                                                                 371 AAGAGGAACGGAGTCCCCGCGCGCGCGCGATTCCCTGAGCTGTGGGACGTGCACC 430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ;
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0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                6.9%; Score 31; DB
100.0%; Pred. No. 13;
cive 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TTGTCTAACCCTAACTGAGAAGGCCGTAGGC 1
SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 84 LENGTH: 79
                                                                                                                                                                                                                                                                                                                                                            431 CAGGACTCGGCTCACACAT 449
                                                                                                                                                                                                                                                                                                                                                                                                               61 caddacrcdcrcacar 79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity 100.
Matches 31, Conservative
                                                                                             ; ORGANISM: Homo sapiens
US-10-714-195-84
                                                                                                                                             Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-330-872-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        -10-330-872-1/c
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LENGTH: 31
                                                                         TYPE: DNA
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Query Match
6.7%; Score 30; DB :
Best Local Similarity 100.0%; Pred. No. 16;
Matches 30; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE DESCRIPTION: SEQ ID NO: 22:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            6.7%; Score 30;
                                                                                                                                                                                                                                                                                                      NAME: Storella, John R. REGISTRATION NUMBER: 32,944
                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 22:
SEQUENCE CHARACTERISTICS:
LENGTH: 30 base pairs
                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION:
                                                       FILING DATE: 07-Feb-2003
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 4, Application US/10330872
Publication No. US20030186282A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Geron Corporation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear MOLECULE TYPE: DNA
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ORGANISM: Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-10-359-935-22
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APPLICANT: Leteinger, Robert L.
APPLICANT: Garimella, Viswanadham
TITLE OF INVENTION: Method of Detection by Enhancement of Silver Staining
FILE REFERENCE: 00-1080 NUMBER: US/09/903,461
CURRENT APPLICATION NUMBER: US/09/903,461
PRIOR APPLICATION NUMBER: 60/217,782
PRIOR PILING DATE: 2000-07-11
PRIOR FILING DATE: 2000-07-11
SOFTWARE: Microsoft Word 98
                                                                                                                                                                                                                                                Gaps
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16;
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STREET: Two Embarcadero Center, Eighth Floor
SITY: San Francisco
STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Feng, Junli
Funk, Walter
Andrewe, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                      6.7%; Score 30; DB 100.0%; Pred. No. 16; ive 0; Mismatches
                                                                                                                                                                                                                                                                                         77 IGCTTTTGCTCCCCGCGCGCTGTTTTCTC 106
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TTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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100.0%; Pred. No.
tive 0; Mismatc
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Publication No. US20030153076A1
GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 2, Application US/09903461
Publication No. US20020034756A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA ORGANISM: Artificial Sequence
TELEFAX: (415) 576-0300
INFORMATION FOR SEG ID NO: 3
SEQUENCE CHARACTERISTICS:
LENGTH: 30 base pairs
                                                                                                                                                                                                    Query Match
Best Local Similarity 100.
Matches 30; Conservative
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                                                                              TYPE: nucleic acid
STRANDEDNESS: sing
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Best Local Similarity
Marches 30; Conserve
                                                                                                                       TOPOLOGY: linear MOLECULE TYPE: DNA US-09-057-351-22
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LENGTH: 30
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APPLICANT: Weinrich, Scott
APPLICANT: Weinrich, Scott
APPLICANT: Atkinson III, Edward
APPLICANT: Lichtstainer, Serge
APPLICANT: Lichtstainer, Serge
APPLICANT: Lichtstainer, Serge
APPLICANT: Pruzan, Ronald
TITLE OF INVENTION: Unibitors
FILE REFERENCE: 011/006C
CURRENT FILING DATE: 2002-12-24
PRIOR APPLICATION NUMBER: 08/510,736
PRIOR APPLICATION NUMBER: 08/833,377
PRIOR APPLICATION NUMBER: 08/833,377
PRIOR PILING DATE: 1997-04-04
PRIOR PILING DATE: 1999-10-18
PRIOR PILING DATE: 1999-10-18
PRIOR FILING DATE: 1999-10-18
PRIOR FILING DATE: 1999-10-18
PRIOR FILING DATE: 1999-10-18
PRIOR FILING DATE: 1090-11-20
NUMBER OF SEQ ID NOS: 11
SOFTWARE: Patentin version 3.1
SEQ ID NO 4

"LENGTH: 30
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SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/359,935
                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: 015389-000821US
                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/057,351
FILING DATE: 08-APPL1994
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-UUL-1994
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-0CT-1994
APPLICATION NUMBER: US 08/472,802
FILING DATE: 07-UUN-1995
ATTORNEY/AGENT INFORMATION:
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Sequence 5, Application US/10811033
| Sequence 5, Application US/10811033
| Publication No. US200500898331
| GENERAL INFORMATION:
| APPLICANT: Geron Corporation
| APPLICANT: Weinrich Scott
| APPLICANT: Atkinson III, Edward
| APPLICANT: Useserot, Alain
| APPLICANT: Useserot, Alain
| APPLICANT: Useserot, Alain
| APPLICANT: Useserot, Alain
| APPLICANT: Delay Puritied Telomerase to Identify Telomerase Activators and
| TITLE OF INVENTION: Inhibitors
| TITLE OF INVENTION: Unhibitors
| TITLE OF INVENTION: Using Puritied Telomerase Activators and
| TITLE OF INVENTION: Using Puritied Telomerase Activators and
| TITLE OF INVENTION: Using Puritied Telomerase (US/10/811,033)
| TITLE OF INVENTION NUMBER: US/10/811,033
| TITLE OF INVENTION NUMBER: US/10/330,872A
| PRIOR APPLICATION NUMBER: 08/610,736
| PRIOR PILING DATE: 1999-10-18
| PRIOR APPLICATION NUMBER: 09/420,056
| PRIOR APPLICATION NUMBER: 09/420,056
| PRIOR APPLICATION NUMBER: 09/420,056
| PRIOR APPLICATION NUMBER: 09/117,828
| PRIOR APPLICATION NUMBER: 09/117,828
| PRIOR APPLICATION NUMBER: 09/10-18
| PRIOR APPLICATION NUMBER: 09/117,828
| PRIOR APPLICATION NUMBER: 09/10-18
| PRIOR PILING DATE: 1999-10-18
| PRIOR PILING DATE: 2000-11-20
| NUMBER OF SEQ ID NOS: 11
| SEQ ID NO 5
| LENGTH: NUMBER: US/10/818
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                                                                                                                                                                                                                                                                                                                                         167 AAACAAAAATGTCAGCTGCTGGCCCGTTC 196
                                                                                                                                                                                                                                                Query Match 6.7%; Score 30; DB Best Local Similarity 100.0%; Pred. No. 16; Matches 30; Conservative 0; Mismatches
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                   PRIOR APPLICATION NUMBER: 09/717,828
PRIOR FILING DATE: 2000-11-20
NUMBER OF SEQ ID NOS: 11
SOFTWARE: PatentIn version 3.1
SEQ ID NO 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; Sequence 3, Application US/10330872; Publication No. US20030186282A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Geron Corporation
APPLICANT: Weinrich, Scott
APPLICANT: Atkinson III, Edward
APPLICANT: Lichtsteiner, Serge
APPLICANT: Vasserot, Alain
APPLICANT: Pruzan, Ronald
                                                                                                                                                      ; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-811-033-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-811-033-5
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US-10-330-972-5/C

i Sequence 5, Application US/10330872

i Publication Wo. US20030186282A1

i GENERAL INFORMATION:

APPLICANT: Geron Corporation

APPLICANT: Weinrich, Scott

APPLICANT: Lichtsteiner, Serge

APPLICANT: Lichtsteiner, Serge

APPLICANT: Lichtsteiner, Serge

APPLICANT: Lichtsteiner, Serge

APPLICANT: Using Purified Telomerase to Identify Telomerase Activators and

ITILE OF INVENTION: Using Purified Telomerase to Identify Telomerase Activators and

ITILE OF INVENTION: Unibitors

FILE REFERENCE: 011/006C

CURRENT APPLICATION NUMBER: 08/510,736

PRIOR APPLICATION NUMBER: 08/510,736

PRIOR FILING DATE: 1999-04-04

PRIOR FILING DATE: 1999-04-04

PRIOR FILING DATE: 1999-01-16

PRIOR PRILING DATE: 1999-01-16

PRIOR PRILING DATE: 1999-10-16

NUMBER OF SEQ ID NOS: 11

SEQ ID NO 5

LENGTH: 30

LENGTH: 30

LENGTH: 30
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                                      Indels
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100.0%; Pred. No. 16;
tive 0; Mismatches 0; Indels
                                      0
                                                                            167 AAACAAAAATGTCAGCTGCCCGTTC 196
100.0%; Pred. No.
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                               30; Conservative
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Best Local Similarity 100.
Matches 30; Conservative
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US-10-330-872-5
         Best Local Similarity
Matches 30; Conserv
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CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/057,351
FILING DATE: 08-APR-1994
CLASSITEATION: 435
PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/08/272,102
FILING DATE: 07-JUL-1994
PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/08/30,123
FILING DATE: 07-JUL-1994
PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/08/472,802
FILING DATE: 07-JUN-1995
ATTORNEY/AGENT INFORMATION:

NAME: Storella, John R.

REFERENCE/POCKET NUMBER: 015389-000821US
TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION SEQ ID NO: 23:

SEQUIRENE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                            ADDRESSEE: Townsend and Townsend and Crew LLP STREEF: Two Embarcadero Center, Eighth Floor CITY: San Francisco STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 30, Application US/09057351
Patent No. US20010034439A1
GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Juni
APPLICANT: Feng, Juni
APPLICANT: Fank, Walter
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
                                           Sequence 23, Application US/09057351
Patent No. US2001003439A1
PAPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junii
APPLICANT: Feng, Junii
APPLICANT: Funk, Walter
APPLICANT: Runk, Walter
APPLICANT: Madrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 5.8%; Score 26; DB Best Local Similarity 100.0%; Pred. No. 30; Matches 26; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS
                                                                                                                                                                                                                                                                                                                                                                                                                  ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: nucleic acid
STRANDEDNESS: sing
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                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY:
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US-20-20-21-1-103-3/6

Publication No. US20050089883A1

GENERAL INROGRATION:

APPLICANT: Weinrich, Scott

APPLICANT: Weinrich, Scott

APPLICANT: Lichtereiner, Serge

APPLICANT: Useriner, Serge

APPLICANT: Useriner, Serge

APPLICANT: NURWINON: Ushing Purified Telomerase to Identify Telomerase Activators and

TITLE OF INVENTION: Ushing Purified Telomerase to Identify Telomerase Activators us applicant APPLICATION NUMBER: US/10/810, 33

FILE REFERENCE: 011/006C

CURRENT APPLICATION NUMBER: US/10/330, 872A

PRIOR FILING DATE: 1995-06-04

PRIOR FILING DATE: 1999-10-18

SEQ ID NOS: 11

SEQ ID NOS: 11

LENGTH: 30
TITLE OF INVENTION: Using Purified Telomerase to Identify Telomerase Activators and TITLE OF INVENTION: Inhibitors
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        412 GAGCTGTGGGACGTGCACCCAGGACTCGGC 441
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 412 GAGCTGTGGGACGTGCACCCAGGACTCGGC 441
                                      THIE REPERENCE: 011/006C
CURRENT APPLICATION NUMBER: US/10/330,872
CURRENT FILING DATE: 2002-12-24
PRIOR APPLICATION NUMBER: 08/510,736
PRIOR FILING DATE: 1995-08-04
PRIOR PILING DATE: 1995-04-04
PRIOR PILING DATE: 1997-04-04
PRIOR PILING DATE: 1999-10-18
PRIOR PILING DATE: 1999-10-18
PRIOR PILING DATE: 2000-11-20
NUMBER OF SEQ ID NOS: 11
SOFTWARE: PRECETIN VERSION 3:1
SOFTWARE: PRECETIN VERSION 3:1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30 GAGCTATGGGACGTGCACCCAGGACTCGGC 1
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Pred. No. 23
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Best Local Similarity 96.7%;
Matches 29; Conservative
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-811-033-3
                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA ORGANISM: Homo Bapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
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CUNDIVAX: USA
CUNDIVAX: USA
CUNDIVAX: USA
CUNDIVER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/NS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/895,606
FILING DATE: 29-Jun-2001
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/710,249
FILING DATE: CUNROWN-
APPLICATION NUMBER: US 60/003,492
FILING DATE: 08-SEP-1995
ATTORNEY/AGENT INFORMATION:
NAME: Stoorella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-001220US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5.8%; Score 26; DB 1; Length 26; 100.0%; Pred. No. 30; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
ADDRESSE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                               CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/895,606
FILING DATE: 29-0un-2001
CLASSIFICATION: «Unknown»
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/710,249
FILING DATE: «Unknown»
APPLICATION NUMBER: 08 60/003,492
FILING DATE: 08-SEP-1995
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-001220US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: DNA
SEQUENCE DESCRIPTION: SEQ ID NO: 25:
US-09-895-606-25
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Publication No. US20030207404A1
GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 25:
SEQUENCE CHARACTERISTICS:
LENGTH: 26 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 100.0
Matches 26; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear
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US-09-895-606-26/c
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Peng, Juniiam H.
Adams, Robert R.
TITLE OF INVENTION: Methods and Reagents for Regulating
Telomere Length and Telomerase Activity
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100.0%; Pred. No. 30;
live 0; Mismatches 0; Indels
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ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSE: Twomsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco
                                                                                                                                                                                                                                                                                        CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/057,351
FILING DATE: 08-APR-1994
CLASSIFICATION NUMBER: US/09/057,351
FILING APPLICATION: 435
PRIOR APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
PRIOR APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION NUMBER: US 08/330,123
FILING DATE: 37-OCT-1994
PRIOR APPLICATION NUMBER: US 08/472,802
FILING DATE: US 08/472,802
FILING DATE: US 08/472,802
FILING DATE: US 08/472,802
FILING DATE: US 08/472,802
STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco STATE: California COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Storella, John R.
REGISSTRATION NUMBER: 32,944
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  45 TCTAACCCTAACTGAGAAGGGCGTAG 70
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Publication No. US20030207404A1
GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
                                                                                                                                                                                                                                                 COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE DOCKET NUMBER: 0153
TELECOMMUNICATION INFORMATION:
TELEPRONE: (415) 576-0200
TELERAX: (415) 576-0200
INFORMATION FOR SEQ ID NO: 30: SEQUENCE CHARACTERISTICS: LENGTH: 26 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                     ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STATE: California
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-09-057-351-30
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Matches 2
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Gaps
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APPLICANT: Cech, Thomas R.
Lingner, Joachim
Nakamura, Toru
Chapman, Karen B.
Morin, Gregg B.
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                                         TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO: 312
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LENGTH: 26 base pai
SEQUENCE CHARACTERISTICS
                                                                                                                                                                                             Query Match
Best Local Similarity 100.0
Matches 26; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                     US-10-044-692-312/c
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Chapman, Karen B.
Morin, Gregg B.
Harley, Calvin
Andrews, William H.
TITLE OF INVENTION: HUMAN THERAPEUTIC METHODS
THERAPEUTIC METHODS
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                                                                                                                                                                                                                                                                                                               Gaps
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MEDIUM TYPE: Floppy disk
COMPUTER: BID PC compatible
COMPUTER: IS PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/044,692
FILING DATE: 11-Jan-2002
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                            5.8%; Score 26; DB 1; Length 26;
100.0%; Pred. No. 30;
Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 335
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET: Two Embarcadero Center, 8th Floor
CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Apple, Randolph T.
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002600US
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/912,951
FILING DATE: «Unkown»
APPLICATION NUMBER: US 08/854,050
FILING DATE: 09-MAY-1997
APPLICATION NUMBER: US 08/851,843
FILING DATE: 06-MAY-1997
APPLICATION NUMBER: US 08/846,017
FILING DATE: 25-APR-1997
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/844,419
FILING DATE: 01-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: United States of America
                                                                                                                                                 TOPOLOGY: linear
MOLECULE TYPE: DNA
SEQUENCE DESCRIPTION: SEQ ID NO: 26:
                                                                                                                                                                                                                                                                                                                                                     145 CTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                                                                                                                                                   26 CTTCCACCGTTCATTCTAGAGCAAAC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 311, Application US/10044692
Publication No. US20030096344A1
GENERAL INFORMATION:
APPLICANT: Cech, Thomas R.
Lingner, Joachim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 311:
  (415) 576-0200
                    (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
                                      INFORMATION FOR SEQ ID NO: 26:
SEQUENCE CHARACTERISTICS:
LENGTH: 26 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                            26; Conservative
  TELEPHONE:
                                                                                                                                                                                                                                                                                 Best Local Similarity
Matches 26; Conserve
                      TELEFAX:
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                                                                                                                                                                                                                                                                 Query Match
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DIAGNOSTIC AND
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COMPUTER: IBM PC compatible
CERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: Twomsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, 8th Floor
CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Apple, Randolph T.
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002600US
                                                                                              DB 1;
30;
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APPLICATION NUMBER: US/10/044,692
FILING DATE: 11-Jan-2002
CLASSIFICATION: «Unknown»
PRIOR APPLICATION: «Unknown»
PRIOR APPLICATION NUMBER: 08/912,951
FILING DATE: «Unknown»
APPLICATION NUMBER: US 08/854,050
FILING DATE: 09-MAY-1997
APPLICATION NUMBER: US 08/854,050
FILING DATE: 06-MAY-1997
APPLICATION NUMBER: US 08/846,017
FILING DATE: 25-APR-1997
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/844,419
FILING DATE: 11-APR-1997
APPLICATION NUMBER: US 08/844,419
FILING DATE: 11-APR-1997
APPLICATION NUMBER: US 08/724,643
                                                                                              5.8%; Score 26; DB 100.0%; Pred. No. 30; tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: United States of America ZIP: 94111
COMPUTER READABLE FORM:
MOLECULE TYPE: DNA
SEQUENCE DESCRIPTION: SEQ ID NO: 311:
US-10-044-692-311
                                                                                                                                                                                               45 TCTAACCCTAACTGAGAGGGCGTAG 70
                                                                                                                                                                                                                                             1 rcraacccraacreagaaggggrag 26
                                                                                                                                                                                                                                                                                                                                                                     Sequence 312, Application US/10044692; Publication No. US20030096344A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION:
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5.8%; Score 26; DB ilarity 100.0%; Pred. No. 30; Conservative 0; Mismatches
  SEQUENCE DESCRIPTION: SEQ ID NO: 311:
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TOPOLOGY: linear
;
MOLECULE TYPE: DNA
; SEQUENCE DESCRIPTION: SEQ 1D NO: 312:
US-10-044-539-312
                                                                                                                                                    45 TCTAACCCTAACTGAGAGGGCGTAG 70
                                                                                                                                                                                          1 rcraacccraacrgagaagggcgrag 26
                                                                                                                                                                                                                                                     RESULT 20
US-10-044-539-312/c
US-10-044-539-312/c
; Sequence 312, Application US/10044539
; Publication No. US20030100093A1
; GENERAL INFORMATION:
; APPLICANT: Cech, Thomas R.
; Infigurer, Joachim Nakamura, Toru
; Chapman, Karen B.
Morin, Geregg B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: (415) 576-0300 INFORMATION FOR SEQ ID NO: 312: SEQUENCE CHARACTERISTICS:
                                           Query Match
Best Local Similarity
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US-10-044-539-311
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TITLE OF INVENTION: HUMAN TELOMERASE CATALYTIC SUBUNIT: DIAGNOSTIC AND THERAPEUTIC METHODS
                                                                                                                                                                      ;
0
                                                                                                                                                                      Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:

REDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/044,539
FILING DATE: 11.Jan-2002
CLASSIFICATION: 435
                                                                                                                          Length 26;
                                                                                                                                                                      0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, 8th Floor CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Apple, Randolph T.
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002600US
                                                                                                                          DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: «UDKNOWN»
APPLICATION NUMBER: US 08/854,050
FILING DATE: 09-MX-1997
APPLICATION NUMBER: US 08/851,843
FILING DATE: 06-MX-1997
APPLICATION NUMBER: US 08/846,017
FILING DATE: 25-APR-1997
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/844,419
FILING DATE: 01-OCT-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                        5.8%; Score 26; DB 100.0%; Pred. No. 30; Live 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: United States of America
                   TOPOLOGY: linear

MOLECULE TYPE: DNA
SEQUENCE DESCRIPTION: SEQ ID NO: 312:
US-10-044-692-312
                                                                                                                                                                                                           145 CTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/912,951
                                                                                                                                                                                                                                        26 CTTCCACCGTTCATTCTAGAGCAAAC 1
                                                                                                                                                                                                                                                                                                                                                       Sequence 311, Application US/10044539
Publication No. USZ0030100093A1
GENERAL INFORMATION:
APPLICANT: Cech, Thomas R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                      Lingner, Joachim
Nakamura, Toru
Chapman, Karen B.
Morin, Gregg B.
Harley, Calvin
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 335
                                                                                                                                            Best Local Similarity 100.
Matches 26; Conservative
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US-10-044-539-311
                                                                                                                          Query Match
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DIAGNOSTIC AND
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Harley, William H.
Andrews, William H.
TITLE OF INVENTION: HUMAN TELOMERASE CATALYTIC SUBUNIT:
THERAPEUTIC METHODS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTER: United States of America

ZIP: 94111
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/044,539
FILING DATE: 11-04n-2002
CLASSIFICATION NUMBER: 08/912,951
FILING DATE: CURNOWN-
APPLICATION NUMBER: 08/912,951
FILING DATE: 06-MAY-1997
APPLICATION NUMBER: US 08/854,050
FILING DATE: 06-MAY-1997
APPLICATION NUMBER: US 08/846,017
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/444,419
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/44,419
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/724,643
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/724,643
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/724,643
FILING DATE: 18-APR-1997
ATTORNEY/AGENT INFORMATION:
NAMAF: ANALS 
     Length 26;
                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 335
CORRESPONDENCE ADDRESS:
ADDRESSE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, 8th Floor
CITY: San Francisco
STATE: California
COUNTRY: United States of America
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Apple, Randolph T.
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002600US
TELECOMMUNICATION INFORMATION:
DB 1;
30;
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Sequence 30, Application US/10359935
Publication No. US20030153076A1
GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 597, Application US/10325810; Publication NO. US2030204069A1; GENERAL INFORMATION: APPLICANT: Cech, Thomas R. Lingner, Joachim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: (415) 576-0300 INFORMATION FOR SEQ ID NO: 30:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: linear
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                  Score 26; DB 1; Length 26; Pred. No. 30;
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100.0%; Pred. No. 30;
                                                          0; Indels
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COMPUTER: IBM PC_COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION UMBER: US/10/359,935
FILING DATE: 07-Feb-2003
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Townsend and Townsend and Crew LL STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-000821US
TELECOMMUNICATION INFORMATION:
TELECHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                          Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/057,351
FILING DATE: 08-APR-1994
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-UUL-1994
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
APPLICATION NUMBER: US 08/32,802
FILING DATE: 07-UUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; Pred. No.
                  5.8%; Score 26; DB
ilarity 100.0%; Pred. No. 30,
Conservative 0; Mismatches
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MOLECULE TYPE: DNA
SEQUENCE DESCRIPTION: SEQ ID NO: 23:
US-10-359-935-23
                                                                                                    145 CTTCCACCGTTCATTCTAGAGCAAAC 170
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                                                                                                                            CTTCCACCGTTCATTCTAGAGCAAAC 1
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                                                                                                                                                                                                                  US-10-359-935-23/c
Sequence 23, Application US/10359935
Publication No. US20030153076A1
GENERAL INFORMATION:
APPLICANT: VIlleponteau, Bryant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 26 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: 23
                                                                                                                                                                                                                                                                                                                         Feng, Junli
Funk, Walter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             26; Conservative
Query Match
Best Local Similarity
Thes 26; Conserve
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Best Local Similarity
Matches 26; Conserva
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US-10-359-935-30
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Chapman, Karen B.
Morin, Gregg B.
Harley, Calvin B.
Andrews, William H.
TITLE OF INVENTION: Human Telomerase Catalytic Subunit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 26;
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                                                                                                                                                                                                                                                                                                                                                                                                                                Version #1.30
                                                                                                                                                           ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco CTATE: California COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-000821US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1;
                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE Floppy disk
COMPUTER READABLE FORM:
MEDIUM TYPE Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PASTENTIN Release #1.0, Ver.
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/359,935
FILING DATE: 07-Feb-2003
CLASSIFICATION: 435
PRIOR APPLICATION NUMBER: US/09/057,351
FILING DATE: 08-APR-1994
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
APPLICATION NUMBER: US 08/330,123
FILING DATE: 07-DCT-1994
APPLICATION NUMBER: US 08/372,102
FILING DATE: 07-DUL-1994
APPLICATION NUMBER: US 08/372,102
FILING DATE: 07-JUL-1994
APPLICATION NUMBER: US 08/472,802
FILING DATE: 07-JUL-1995
Feng, Junli
Funk, Walter
Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 5.8%; Score 26; DB Best Local Similarity 100.0%; Pred. No. 30; Matches 26; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; SEQUENCE DESCRIPTION: SEQ ID NO: 30:
US-10-359-935-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               45 TCTAACCCTAACTGAGAAGGGCGTAG 70
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Morin, Gregg B.
Harley, Calvin B.
Andrews, William H.
TITLE OF INVENTION' Human Telomerase Catalytic Subunit
NUMBER OF SEQUENCES: 633
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARRE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/325,810
FILING DATE: 20-Dec-2002
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                            ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION: CUNknown>
PRIOR APPLICATION DATA:

APPLICATION DATA:

FILING DATE: 29-Sep-1997

APPLICATION NUMBER: US 08/724,643

FILING DATE: 01-CCT-1996

APPLICATION NUMBER: US 08/844,419

FILING DATE: 18-APR-1997

APPLICATION NUMBER: US 08/844,419

FILING DATE: 18-APR-1997

APPLICATION NUMBER: US 08/851,843

FILING DATE: 25-APR-1997

APPLICATION NUMBER: US 08/851,843

FILING DATE: 06-MAY-1997

APPLICATION NUMBER: US 08/851,550

FILING DATE: 14-AUG-1997

APPLICATION NUMBER: US 08/912,951

FILING DATE: 14-AUG-1997

APPLICATION NUMBER: US 08/912,951

FILING DATE: 14-AUG-1997

APPLICATION NUMBER: US 08/915,503

FILING DATE: 14-AUG-1997

APPLICATION NUMBER: US 08/915,503

FILING DATE: 14-AUG-1997

APPLICATION NUMBER: US 08/915,503

FILING DATE: 11-AUG-1997

**TORREY/AGENT INFORMATION**
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Ausenhus, Scott L.
REGISTRATION NUMBER: 42,271
REFERENCE/DOCKET NUMBER: 015389-002620US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; SEQUENCE DESCRIPTION: SEQ ID NO: 598:
US-10-325-810-598
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              145 CTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
                                                                     Nakamura, Toru
Chapman, Karen B.
                                      Lingner, Joachim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (415) 576-0300
                APPLICANT: Cech, Thomas R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 26 base pairs
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                                                                                                                                                                                                                                                                                                                                   CITY: San Francisco
STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: -
LOCATION: 1..26
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE: DNA
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0
                                                                                                                                                                                                                                                                                    CURROLLER: LEW PC COMBALIDATE
CURROLLER: PATENT: PC-DOS/MS-DOS
SOFTWARE: PATENTING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTING DATA:
APPLICATION NUMBER: US/10/325,810
FILING DATE: 20-Dec-2002
CLASSIFTCATION: «Unknown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/402,181
FILING DATE: 29-Sep-1997
APPLICATION NUMBER: US 08/844,419
FILING DATE: 01-OCT-1996
APPLICATION NUMBER: US 08/844,419
FILING DATE: 04-APR-1997
APPLICATION NUMBER: US 08/846,017
FILING DATE: 06-MAY-1997
APPLICATION NUMBER: US 08/854,050
FILING DATE: 09-MAY-1997
APPLICATION NUMBER: US 08/854,050
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NUMBER OF SEQUENCES: 613
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Ausenhus, Scott L.
REGISTRATION NUMBER: 42,271
REFERENCE/DOCKET NUMBER: 015389-002620US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            .;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/912,951
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/915,503
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: WO PCT/US97/17885
FILING DATE: 01-OCT-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 5.8%; Score 26; DB 1; Best Local Similarity 100.0%; Pred. No. 30; Matches 26; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TCTAACCCTAACTGAGAAGGGCGTAG 70
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                                                                                                                                                                                         ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: F10Ppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 24
US-10-325-810-598/c
; Sequence 599, Application US/10325810
; Publication No. US20030204069A1
; CENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 26 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO: 597
                                                                                                              CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY: -
LOCATION: 1..26
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY:
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US-10-877-124-597
Sequence 597, Application US/10877124
Publication No. US20040242529A1
Seneral INFORMATION:
Information (Cech, Thomas R.
Information (Chapman, Toru Chapman, Karen B.
Morin, Greeg B.
Harley, Calvin B.
Andrews, William H.
TITLE OF INVENTION: Human Telomerase Catalytic Subunit NUMBER OF SEQUENCES: 727
CORRESPONDENCE ADDRESS:
STRUKE TWO Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTY: CALLLUILLE
SIAIR: CALLLUILLE
COUNTYER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENT: PC-DOS/MS-DOS
SOFTWARE: PATENT: Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
PILING DATE: 24-Jun-2004
CLASSIFICATION NUMBER: US/10/877,124
FILING DATE: CANNOWN-1999
APPLICATION NUMBER: US/999
APPLICATION NUMBER: US/999
APPLICATION NUMBER: US/974,549
FILING DATE: CHRICOWN-1997
APPLICATION NUMBER: US/974,549
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US/97
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REGISTRATION NUMBER: 36,429
REFRENCE/DOCKET NUMBER: 015389-002610US
TELECOWMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NOMBER: US 09/031/030

FILING DATE: 09-MAY-1997

APPLICATION NUMBER: US 08/911,312

FILING DATE: 14-AUG-1997

APPLICATION NUMBER: US 08/912,951

FILING DATE: 14-AUG-1997

APPLICATION NUMBER: WO PCT/US97/17618

FILING DATE: 01-0CT-1997

APPLICATION NUMBER: WO PCT/US97/17618

PILING DATE: 01-0CT-1997

APPLICATION NUMBER: WO PCT/US97/17685

PILING DATE: 01-0CT-1997

ATTORNEY/AGENT INFORMATION:
26 CTTCCACCGTTCATTCTAGAGCAAAC 1
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LENGTH: 26 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO: 597
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear
MOLECULE TYPE: DNA
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LOCATION: 1..26
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Publication No. US20040242529A1

GENERAL INFORMATION:
Lingner, Joachim
Nakamura, Toru
Chapman, Karen B.
Morin, Gregg B.
Harley, Calvin B.
Andrews, William H.
TITLE OF INVENTION: Human Telomerase Catalytic Subunit
NUMBER OF SEQUENCES: 727

CORRESPONDENCE ADRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:

MEDLUM TYPE: Floppy disk

MEDLUM TYPE: Floppy disk

MEDLUM TYPE: Floppy disk

MEDLUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/10/877,124

FILING DATE: 24-Jun-2004

FILING DATE: COLNOW-1999

APPLICATION NUMBER: US/99/4,549

FILING DATE: COLNOW-1997

APPLICATION NUMBER: US 08/844,419

FILING DATE: 25-APR-1997

APPLICATION NUMBER: US 08/851,843

FILING DATE: 25-APR-1997

APPLICATION NUMBER: US 08/851,843

FILING DATE: 06-MAY-1997

APPLICATION NUMBER: US 08/851,312

APPLICATION NUMBER: US 08/911,312

APPLICATION NUMBER: US 08/911,312

FILING DATE: 14-AUG-1997

APPLICATION NUMBER: US 08/912,951

FILING DATE: 14-AUG-1997

APPLICATION NUMBER: US 08/912,951

FILING DATE: 14-AUG-1997

APPLICATION NUMBER: US 08/912,951
                                                                                                                Length 26;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco CATTE: California COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Apple, Randolph Ted
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002610US
; SEQUENCE DESCRIPTION: /note= "F3b primer"; SEQUENCE DESCRIPTION: SEQ ID NO: 597: US-10-877-124-597
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: WO PCT/US97/17618
FILING DATE: 01-0CT-1997
APPLICATION NUMBER: WO PCT/US97/17885
FILING DATE: 01-0CT-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                DB 1;
30;
                                                                                                              Query Match 5.8%; Score 26; DB Best Local Similarity 100.0%; Pred. No. 30; Matches 26; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US 08/915,503
FILING DATE: 14-AUG-1997
                                                                                                                                                                                                                      45 TCTAACCCTAACTGAGAAGGGCGTAG 70
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TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                            US-10-877-124-598/c
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Score 26; DB 1;
Pred. No. 30;
0; Mismatches
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                                                                                                                                                            TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0200
INFORMATION FOR SEG ID NO: 597:
SEQUENCE CHARACTERISTICS:
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100.0%; Pre
0; !
                                                                                                                                                                                                                                      LENGTH: 26 base pairs
                                                                                                                                                                                                                                                          TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                TOPOLOGY: linear
MOLECULE TYPE: DNA
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 100.
Matches 26; Conservative
                                                                                                                                                                                                                                                                                                                                                        NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-10-877-022-598/c
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Publication No. US20040247613A1

GENERAL INFORMATION:

Lingner, Joachim

Nakamura, Toru

Chapman, Karen B.

Morin, Gregg B.

Harley, Calvin B.

Andrews, William H.

TITLE OF INVERTYION: Human Telomerase Catalytic Subunit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUPTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMFUTER: IBM PC compatible
COMFUTER: Ploppy disk
COMFUTER: Ploppy disk
COMFUTER: DATA:
COMFUTER: DATA:
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
FILING DATE: 24-Jun-2004
CLASSIFICATION: <UNKNOWN>
                                                                                                                                                                                                                                                                      5.8%; Score 26; DB 1; Length 26; 100.0%; Pred. No. 30; Aiswarches 0; Indels native
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
                                                                                                                              ) NAME/KEY: - LOCATION: 1..26
) COTHER INFORMATION: /note= "R3c primer";
SEQUENCE DESCRIPTION: SEQ ID NO: 598:
US-10-877-124-598
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: WO PCT/US97/17618
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APPLICATION NUMBER: US/09/432,503
FILING DATE: 02-Nov-1999
FILING DATE: 08/974,549
FILING DATE: vUMBER: 08/974,549
APPLICATION NUMBER: US 08/844,419
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/846,017
FILING DATE: 25-APR-1997
APPLICATION NUMBER: US 08/851,843
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: 09-MAY-1997
PAPLICATION NUMBER: US 08/911,312
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/912,951
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US 08/854,050
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/915,503
FILING DATE: 14-AUG-1997
                                                                                                                                                                                                                                                                                                                                                145 CTTCCACCGTTCATTCTAGAGCAAAC 170
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INFORMATION FOR SEQ ID NO: 598:
SEQUENCE CHARACTERISTICS:
LENGTH: 26 base pairs
                                                            TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STATE: California
                                                                                              TOPOLOGY: linear
MOLECULE TYPE: DNA
FEATURE:
                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 27
US-10-877-022-597
                                                                                                                                                                                                                                                                                                            26;
                                                                                                                                                                                                                                                                      Query Match
Best Local
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Gaps
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; Publication No US20040247613A1
; GENERAL INFORMATION:
    Lingmer, Joachim
    Nakamura, Toru
    Chapman, Karen B.
    Maril, Gregg B.
    Harley, Calvin B.
    Andrews, William H.
    TITLE OF INVENTION: Human Telomerase Catalytic Subunit
    NUMBER OF SEQUENCES: 727
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: CALILOINIA
COUNTRY: USA
ZIP: 9411-3834
ZIP: 9411-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/877,022
FILING DATE: 24-Jun-2004
CLASSIFICATION NUMBER: US/99/432,503
FILING DATE: CONOV-1999
APPLICATION NUMBER: US/99/4549
FILING DATE: CUNKNOWN>
APPLICATION NUMBER: US/99/4,549
FILING DATE: CUNKNOWN>
APPLICATION NUMBER: US/98/44,419
FILING DATE: LARAPH-1997
APPLICATION NUMBER: US/98/846,017
FILING DATE: S2-APR-1997
APPLICATION NUMBER: US/98/846,017
FILING DATE: S1-APR-1997
APPLICATION NUMBER: US/98/846,017
FILING DATE: S1-APR-1997
APPLICATION NUMBER: US/98/846,017
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 26;
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ADDRESSE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco STATE: California
APPLICATION NUMBER: WO PCT/US97/17885
FILING DATE: 0.1-0CT-1997
ATTORNEY/AGENT INFORMATION:
NAME: Apple, Randolph Ted
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 36,429
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ;;
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OTHER INFORMATION: /note= "F3b primer"
SEQUENCE DESCRIPTION: SEQ ID NO: 597:
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Name of the control o
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REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002610US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE DESCRIPTION: SEQ ID NO: 597:
US-10-877-146-597
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 14-AUG-1997
APPLICATION WUMBER: US 08/915,503
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: WO PCT/US97/17618
FILING DATE: 01-OCT-1997
APPLICATION NUMBER: WO PCT/US97/17885
FILING DATE: 01-OCT-1997
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   FILING DATE: 02-Nov-1999
APPLICATION NUMBER: 08/974,549
FILING DATE: «Unknown»
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APF-1997
FILING DATE: 25-APR-1997
                                                                                                                                                                                                                                                  APPLICATION NUMBER: US 08/851,843
FILING DATE: 06-MAY-1997
PILING DATE: 09-MAY-1997
FILING DATE: 09-MAY-1997
APPLICATION NUMBER: US 08/911,312
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/912,951
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 5.8%; Score 26; DB Best Local Similarity 100.0%; Pred. No. 30; Matches 26; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      45 TCTAACCCTAACTGAGAAGGGCGTAG 70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 TCTAACCCTAACTGAGAAGGGCGTAG 26
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 26 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: (415) 576-03
INFORMATION FOR SEQ ID NO: 597
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LOCATION: 1..26
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-10-877-146-598/c
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Publication No. US20050013825A1
GENERAL INFORMATION:
Ingrer, Joachim
Nakamura, Toru
Chapman, Karen B.
Morin, Gregg B.
Harley, Calvin B.
Andrews, William H.
TITLE OF INVENTION: Human Telomerase Catalytic Subunit
NUMBER OF SEQUENCES: 727
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 5.8%; Score 26; DB 1; Length 26;
100.0%; Pred. No. 30;
.ive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Apple, Randolph Ted
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002610US
TELECOMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 598:
PILING DATE: 09-MAY-1997

APPLICATION NUMBER: US 08/911,312

FILING DATE: 14-AUG-1997

APPLICATION WUMBER: US 08/912,951

FILING DATE: 14-AUG-1997

APPLICATION WUMBER: US 08/915,503

FILING DATE: 14-AUG-1997

APPLICATION NUMBER: WO PCT/US97/17618

FILING DATE: 01-OCT-1997

APPLICATION NUMBER: WO PCT/US97/17618

FILING DATE: 01-OCT-1997

ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: /note= "R3c primer"
SEQUENCE DESCRIPTION: SEQ ID NO: 598:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/10/877,146
FILING DATE: 24-Jun-2004
CLASSIFICATION: «Unknown»
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                145 CTTCCACCGTTCATTCTAGAGCAAAC 170
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 26 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LOCATION: 1..26
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STATE: California
COUNTRY: USA
ZIP: 94111-3834
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity 100.
Matches 26; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 29
US-10-877-146-597
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TITLE OF INVENTION: Mammalian H.

TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Bighth Floor
CITY: San Francisco
STATE: California
COUNTRY: San Francisco
STATE: 4111-3834
COMPUTER: READABLE FORM:
MEDIUM TYPE: Flooppy disk
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        5.5%; Score 25; DB 1; Length 28;
100.0%; Pred. No. 46;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                             COMPUTER READABLE FORM:

MEDIUW TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATURG SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/057,351
FILING DATE: 08-APR-1994
CLASSIFICATION NUMBER: US 08/272,102
FILING DATE: O'-JUL-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: O'-JUL-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/472,802
FILING DATE: O'-JUN-1995
APPLICATION NUMBER: US 08/472,802
FILING DATE: O'-JUN-1995
APPLICATION NUMBER: US 08/472,802
FILING DATE: O'-JUN-1995
APPLICATION NUMBER: US 08/472,802
FILING DATE: 18-DAMATION:
TELEBROME: Storella, John R.
REFERENCY DOCKET NUMBER: 015389-000821US
TELECOMMUNICATION INFORMATION:
TELEBROME: (415) 576-0200
TELEBROME: GRANGTERISTICS:
BEQUENCE CHARACTERISTICS:
BEQUENCE CHARACTERISTICS:
CLENGTH: US BASE PRICE
TYPE: NUCLEIC acid
                         CORRESPONDENCE ADDRESS:
ADDRESSE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Bighth Floor CITY: San Francisco STATE: California COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  81 TTTGCTCCCGCGCGCTGTTTTCT 105
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-10-359-935-29/c
; Sequence 29, Application US/10359935
; Publication No. US20030153076A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Feng, Junli
Funk, Walter
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                                                                                                                                                                                       ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
     NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 linear
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tes 25; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Loc
Matches
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0
COMPUTER READABLE FORM
MEDIUM TYPE: Floppy disk
COMPUTER: IRM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/10/877,146
FILING DATE: 24-Jun-2004
CLASSIFICATION NUMBER: US/09/432,503
FILING DATE: 24-Jun-2004
CLASSIFICATION NUMBER: US/09/432,503
FILING DATE: 02-NOV-1999
FILING DATE: 02-NOV-1999
APPLICATION NUMBER: US 08/844,419
FILING DATE: 12-APR-1997
APPLICATION NUMBER: US 08/854,050
FILING DATE: 12-APR-1997
APPLICATION NUMBER: US 08/854,050
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/81,843
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/912,951
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: WO PCT/US97/17885
FILING DATE: 01-OCT-1997
APPLICATION NUMBER: WO PCT/US97/17885
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Apple, Randolph Ted
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002610US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LOCATION: 1..26

COTHER INFORMATION: /note= "R3c primer"

SEQUENCE DESCRIPTION: SEQ ID NO: 598:
US-10-877-146-598
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Patent No. US200100343391
GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junli
APPLICANT: Funk, Walter
APPLICANT: Marews, William H.
TITLE OF INVENTION: Mammalian Telomerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 145 CTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         26 CTTCCACCGTTCATTCTAGAGCAAAC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 26 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO: 598:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear
MOLECULE TYPE: DNA
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY:
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US-09-057-351-29/c
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RESULT 34

US-09-018-125-5/C
US-09-018-125-5/C

Sequence 5, Application US/09018125A

Patent No. US2001000902A1

GENERAL INFORMATION:
APPLICANT: Sidverman, Robert H.
APPLICANT: Cowell, John K.
APPLICANT: Cowell, John K.
APPLICANT: Torrence, Paul F.
TITLE OF INVENTION: RNASE L ACTIVATORS AND ANTISENSE OLIGONUCLEOTIDES
TITLE OF INVENTION: EFFECTIVE TO TREAT TELOMERASE-EXPRESSING MALIGNANCIES
FILE REFERENCE: 8656-022

CURRENT PILIOR DATE: 1999-02-03

EARLIER APPLICATION NUMBER: 6/044,507

EARLIER FILING DATE: 1999-04-21

WUMBER OF SEQ ID NOS: 9

SOFTWARE: PatentIn Ver. 2.0

LENGTH: 24

TENDTH: 24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 5.3%; Score 24; DB 1; Length 24; Best Local Similarity 100.0%; Pred. No. 41; Matches 24; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
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Fublication No. US20050019785A1
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Braker, Joffre
APPLICANT: Cronin, Maureen
APPLICANT: Grak, Steve
TITLE OF INVENTION: GENE EXPRESSION PROFILING OF EGFR
TITLE OF INVENTION: GENE EXPRESSION PROFILING
FILE REFERENCE: 39740-0005
CURRENT APPLICATION NUMBER: US/10/714,195
CURRENT APPLICATION NUMBER: 60/427090
PRIOR FILING DATE: 2003-11-14
PRIOR APPLICATION NUMBER: 60/427090
PRIOR PRILNG DATE: 2003-11-15
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 345
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 5.1%; Score 23; DB 1; Best Local Similarity 100.0%; Pred. No. 48; Matches 23; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               423 CGTGCACCCAGGACTCGGCTCACA 446
1 TTTGTCTAACCCTAACTGAGAAGG 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     404 GATTCCCTGAGCTGTGGGACGTG 426
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; OTHER INFORMATION: primer US-10-714-195-345
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 35
US-10-714-195-345/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-09-018-125-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          g
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Patent No. US20010007902A1;
GENURAL INFORMATION:
APPLICANT: Silverman, Robert H.
APPLICANT: Cowell, John K.
APPLICANT: Cowell, John K.
APPLICANT: Torrence, Paul F.
TITLE OF INVENTION: RNASE L ACTIVATORS AND ANTISENSE OLIGONUCLEOTIDES
TITLE OF INVENTION: RNASE L ACTIVATORS AND ANTISENSE OLIGONUCLEOTIDES;
TITLE OF INVENTION: BFFECTIVE TO TREAT TELOMERASE-EXPRESSING MALIGNANCIES;
TITLE OF INVENTION: BFFECTIVE TO TREAT TELOMERASE-EXPRESSING MALIGNANCIES;
CURRENT APPLICATION NUMBER: US/09/018,125A;
CURRENT FILING DATE: 1999-02-03;
EARLIER PELING DATE: 1999-04-21;
NUMBER OF SEQ ID NOS: 9
SOFTWARE: PALENTIN Ver. 2.0;
SEQ ID NO 4;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 5.5%; Score 25; DB 1; Length 28; Best Local Similarity 100.0%; Pred. No. 46; Matches 25; Conservative 0; Mismatches 0; Indels
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  SOFTWARE: PatentIn Release #1.0, Version #1.30
                                           PAPELICATION NUMBER: US/10/359,935
FILING DATE: 07-Feb-2003
CLASSIFICATION: 435
PRIOR APPLICATION HOWER: US/09/057,351
FILING DATE: 08-APPLICATION NUMBER: US/09/057,351
FILING DATE: 07-JUL-1994
APPLICATION NUMBER: US 08/30,123
FILING DATE: 27-GT-1994
APPLICATION NUMBER: US 08/30,123
FILING DATE: 07-JUL-1994
APPLICATION NUMBER: US 08/30,123
FILING DATE: 07-GT-1994
APPLICATION NUMBER: US 08/30,123
FILING DATE: 07-GT-1994
APPLICATION NUMBER: US 08/30,123
FILING DATE: 07-GT-1994
APPLICATION NUMBER: US 08/472,802
FILING BATE: 07-GT-1995
ATTORNEY/AGENT INFORMATION:
NAME: SECRETATION NUMBER: 32,944
FREFERENGE/DOCKET NUMBER: 015389-000821US
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: DNA
SEQUENCE DESCRIPTION: SEQ ID NO: 29:
US-10-359-935-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         81 TITGCICCCGCGCGCTGTTTTCT 105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                41 TTTGTCTAACCCTAACTGAGAAGG 64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (415) 576 0300
INFORMATION FOR SEQ ID NO: 29:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 28 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: nucleic acid
STRANDEDNESS: single
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear
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; OTHER INFORMATION: Description of Artificial Sequence: Target Sequence/sinA sense re
US-10-923-330-511
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APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Biselgen, James
APPLICANT: McSwiggen, Jeonid
TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
CURRENT FILING DATE: 2004-08-20
FRIOR PELLORION NUMBER: US (0/396,600
FRIOR FILING DATE: 2002-07-17
FRIOR APPLICATION NUMBER: US (0/396,600
FRIOR PILING DATE: 2004-05-24
FRIOR PELLORION NUMBER: US 10/826,966
FRIOR FILING DATE: 2004-01-14
FRIOR PELLORION NUMBER: US 10/757,803
FRIOR PELLORION NUMBER: US 10/757,803
FRIOR PELLORION NUMBER: US 10/693,059
FRIOR PELLORION NUMBER: US 10/693,059
FRIOR PELLING DATE: 2003-11-23
FRIOR APPLICATION NUMBER: US 10/693,059
FRIOR PELLING DATE: 2003-02-20
FRIOR APPLICATION NUMBER: PCT/US03/0508
FRIOR APPLICATION NUMBER: PCT/U
               PRIOR FILING DATE: 2004-04-16
PRIOR PELING DATE: 2004-04-17
PRIOR PELING DATE: 2004-04-14
PRIOR APPLICATION NUMBER: US 10/757,803
PRIOR FILING DATE: 2003-01-14
PRIOR PELING DATE: 2003-11-24
PRIOR PELING DATE: 2003-11-24
PRIOR PELING DATE: 2003-11-24
PRIOR PELING DATE: 2003-11-23
PRIOR PELING DATE: 2003-05-23
PRIOR APPLICATION NUMBER: US 10/44,853
PRIOR APPLICATION NUMBER: PCT/US03/05346
PRIOR APPLICATION NUMBER: PCT/US03/05346
PRIOR PILING DATE: 2003-02-20
PRIOR FILING DATE: 2003-02-20
PRIOR FILING DATE: 2003-02-20
PRIOR FILING DATE: 2003-03-22
PRIOR PRIOR PREDICATION NUMBER: PCT/US03/05028
PRIOR FILING DATE: 2003-03-23
PRIOR PLING DATE: 2003-03-23
PRIOR FILING DATE: 2003-03-23
PRIOR PLING DATE: 2003-03-23
PRIOR PLING DATE: 2003-02-30
PRIOR FILING DATE: 2003-02-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 5.1%; Score 23; DB 1; Length 23; Best Local Similarity 73.9%; Pred. No. 48; Matches 17; Conservative 6; Mismatches 0; Indels
PRIOR APPLICATION NUMBER: US 10/826,966
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             136 GCCTGCCGCCTTCCACCGTTCAT 158
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 GCCUGCCGCCUUCCACCGUUCAU 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 38
US-10-923-330-512
Sequence 512, Application US/10923330
Publication No. US20050153916A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: RNA
ORGANISM: Artificial Sequence
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LENGTH: 23
TYPE: RNA
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                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2 GGTTGCGGAGGGTGGGCCTGGGA 24
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ORGANISM: Artificial Sequence
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Best Local Similarity
Matches 19; Conserva
         RESULT 36
US-10-923-330-510
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LENGTH: 23
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Gaps

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OTHER INFORMATION: Description of Artificial Sequence: Target Sequence/#inA sense re
                                   Sequence 516, Application US/10923330

Publication No. US20050153916A1

GENERAL INFORMATION:
APPLICANT: Sirna Florapoutics, Inc.
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: RAN INTERference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
FILE REFERENCE: 400/209 (MBHB02-708-C)
FURRENT PELING DATE: 2004-08-20
PRIOR APPLICATION NUMBER: 9C1/0803/04088
PRIOR FILING DATE: 2002-07-17
PRIOR APPLICATION NUMBER: US 60/396,600
PRIOR FILING DATE: 2002-07-17
PRIOR PELING DATE: 2004-05-24
PRIOR PELING DATE: 2004-06-24
PRIOR PELING DATE: 2004-06-14
PRIOR PELING DATE: 2004-01-14
PRIOR FILING DATE: 2004-01-14
PRIOR PRIOR APPLICATION NUMBER: US 10/757,803
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         143 GCCTTCCACCGTTCATTCTAGAG 165
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     McSwiggen, James
Beigelman, Leonid
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                                                     ; OTHER INFORMATION: Description of Artificial Sequence: Target Sequence/sinA sense US-10-923-330-512
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US-10-923-330-513
                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 513, Application US/10923330
| Publication No. US20050153916A1
| GENERAL INFORMATION: US20050153916A1
| GENERAL INFORMATION: US20050153916A1
| GENERAL INFORMATION: US20050153916A1
| APPLICANT: Sirna Therapeutics, Inc.
| APPLICANT: MCSW4Gen, James
| APPLICANT: MCSW4Gen, James
| APPLICANT: BINDAMEDIAN INTERFERENCE: 400/209 | MRH02-708-C)
| TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
| FILE REFERENCE: 400/209 | MRH02-708-C)
| CURRENT FILING DATE: 2004-08-20
| FRICK APPLICATION NUMBER: US 6/396,600
| FRICK APPLICATION NUMBER: US 6/396,600
| FRICK FILING DATE: 2002-07-17
| FRICK FILING DATE: 2003-01-14
| FRICK FILING DATE: 2003-11-23
| FRICK FILING DATE: 2003-12-20
| FRICK FILING DATE: 2003-12-23
| FRICK FILING DATE: 2003-05-23
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                                                                                                                                   Query Match 5.1%; Score 23; DB 1; Length 23; Best Local Similarity 95.7%; Pred. No. 48; Matches 22; Conservative 1; Mismatches 0; Indels
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                                                                                                                                                                                                                                                           283 GCACCCACTGCCACCGCGAAGAG 305
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         395 GCGCGCGCGATTCCCTGAGCTG 417
                                                                                                                                                                                                                                                                                         1 GCACCCACUGCCACCGCGAAGAG 23
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; Sequence 515, Application US/10923330
; Publication No. US20050153916A1
; GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Artificial Sequence
                                                                                                                                         Query Match
Best Local Similarity
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; OTHER INFORMATION: Description of Artificial Sequence: Target Sequence/siNA sense re
US-10-923-330-518
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APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Beigelman, Leonia
APPLICANT: Beigelman, Leonia
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RNA Interference: 1004.029 (MEMBER: US/10/20) 4088

FILE REPERENCE: 400.020 (MEMBER: US/10/20) 4088

PRIOR PELING DATE: 2004-05-24

PRIOR PELING DATE: 2004-05-24

PRIOR PELING DATE: 2004-01-17

PRIOR PELING DATE: 2004-01-17

PRIOR PELING DATE: 2004-01-14

PRIOR PELING DATE: 2004-01-14

PRIOR PELING DATE: 2004-01-14

PRIOR PELING DATE: 2003-01-12-4

PRIOR PELING DATE: 2003-01-12-4

PRIOR PELING DATE: 2003-01-13-4

PRIOR PELING DATE: 2003-01-3-3

PRIOR PELING DATE: 2003-01-3-3

PRIOR PELING DATE: 2003-02-20

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Publication No. US20050153916A1
GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT FILING DATE: 2004-08-20
                                              Gaps
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                                          Indels
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                                          ;
0
                                      7; Mismatches
                                                                                                   145 CTTCCACCGTTCATTCTAGAGCA 167
                                                                                                                                                     1 CUUCCACCGUUCAUUCUAGAGCA 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           146 TTCCACCGTTCATTCTAGAGCAA 168
                                                                                                                                                                                                                                                                                RESULT 43
US-10-923-330-518
US-10-923-330-518
Sequence 518, Application US/10923330
Publication No. US20050153916A1
GENERAL INFORMATION:
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                                      16; Conservative
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                              Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Target Sequence/sinA sense
US-10-923-330-516
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OTHER INFORMATION: Description of Artificial Sequence: Target Sequence/sinA sense
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Sequence 517, Application US/10923330

PUDIcation No. US20050153916A1

GENERAL INFORMATION:

APPLICANT: Sirna Therapeutics, Inc.

APPLICANT: McSwiggen, James

FILE OF INVENTION: Expression Using Short Interfering RNA (sinA)

FILE REFERENCE: 400/209 (MHHBO)-708-C)

FRICK FILING DATE: 2004-08-20

FRIOR PILING DATE: 2004-08-20

FRIOR FILING DATE: 2002-07-17

FRIOR FILING DATE: 2004-01-14

FRIOR FILING DATE: 2004-01-14

FRIOR FILING DATE: 2004-01-14

FRIOR FILING DATE: 2004-01-14

FRIOR FILING DATE: 2003-11-23

FRIOR FILING DATE: 2003-05-23

FRIOR FILING DATE: 2003-05-23

FRIOR FILING DATE: 2003-05-34

FRIOR FILING DATE: 2003-05-34

FRIOR FILING DATE: 2003-05-36

FRIOR FILING DATE: 2003-05
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PRIOR FILING DATE: 2003-11-23
PRIOR APPLICATION NUMBER: US 10/444,853
PRIOR FILING DATE: 2003-05-23
PRIOR FILING DATE: 2003-05-20
PRIOR FILING DATE: 2003-02-20
PRIOR FILING DATE: 2003-02-20
PRIOR FILING DATE: 2003-02-20
Remaining Prior Application data removed - See File Wrapper or PALM, SOFTYARRE: Patentin version 3.3
SOFTYARE: Patentin version 3.3
LENGTH: 23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         5.1%; Score 23; DB 1; Length 23; 69.6%; Pred. No. 48; 0; Indels tive 7; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 23;
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48;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       144 CCTTCCACCGTTCATTCTAGAGC 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: RNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: RNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 69.6%
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 42
US-10-923-330-517
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-10-923-330-517
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; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Target Sequence/sinA sense re
US-10-923-330-520
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US-10-923-330-521
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Remaining Prior Application data removed - See File Wrapper or PALM. NUMBER OF SEQ ID NOS: 768 SOFTWARE: PatentIn version 3.3 SEQ ID NO 520
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
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                                                                                                                                                                                                                                                                                                             1; Length 23;
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                                                                                                                                                                                                                                                                                                       5.1%; Score 23; DB 78.3%; Pred. No. 48; tive 5; Mismatches
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                                                                                                                                                  TYPE: RNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 78.33
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 78.3
Matches 18; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: BLIGHGUILER, INC.
APPLICANT: BASWIGGEN, James
APPLICANT: Beigelman, Leonid
APPLICANT: Beigelman, Leonid
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RAPELSATION WINGER: 400/209 (MBHB02-708-C)
CURRENT PELLORION NUMBER: US/10/923,330
CURRENT PELLORION NUMBER: PCT/US03/0408
PRIOR APPLICATION NUMBER: PCT/US03/0408
PRIOR FILING DATE: 2004-05-24
PRIOR APPLICATION NUMBER: US 60/396,600
PRIOR FILING DATE: 2004-05-24
PRIOR PELLOR DATE: 2004-06-24
PRIOR PELLOR DATE: 2004-01-14
PRIOR PELLOR DATE: 2004-01-14
PRIOR PELLOR DATE: 2003-11-24
PRIOR PELLOR DATE: 2003-11-23
PRIOR PELLOR DATE: 2003-11-23
PRIOR PELLOR DATE: 2003-11-23
PRIOR PELLOR DATE: 2003-11-24
PRIOR PELLOR DATE: 2003-11-23
PRIOR PELLOR DATE: 2003-11-23
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  PRIOR APPLICATION NUMBER: PCT/US03/04088
PRIOR FILING DATE: 2004-05-24
PRIOR APPLICATION NUMBER: US 60/396,600
PRIOR APPLICATION NUMBER: US 60/396,600
PRIOR PILING DATE: 2004-05-24
PRIOR PLING DATE: 2004-05-24
PRIOR PLING DATE: 2004-01-6
PRIOR PLING DATE: 2004-01-14
PRIOR FILING DATE: 2004-01-14
PRIOR PLING DATE: 2003-11-24
PRIOR APPLICATION NUMBER: US 10/720,448
PRIOR PLING DATE: 2003-11-24
PRIOR PLING DATE: 2003-02-20
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APPLICATION NUMBER: PCT/US03/05028
FILING DATE: 2003-02-20
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Publication No. US20050153916A1
GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
APPLICANT: Beigelman, Leonid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: RNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-10-923-330-519
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LENGTH: 23
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4.9%; Score 22; DB 1; Length 22; 100.0%; Pred. No. 55; 0; Indels ive 0; Mismatches 0; Indels
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Prancisco
                                                                                                                   ZIP: 9411.3834

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPATER: BATENING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/057,351
FILING DATE:
E: Townsend and Townsend and Crew LLP
Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION: 43.5
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-UUL-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/472,802
FILING DATE: 07-UN-1995
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R:
REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944
REGISTRATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION 118FORMATION:
TELEFAX: (415) 576-0300
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Funk, Walter
Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 41, Application US/10359935
Publication No. US20030153076A1
GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     22 AACTGAGAAGGCGTAGGCGCC 1
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COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     54 AACTGAGAAGGGCGTAGGCGCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 42:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 22 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 100.
Matches 22; Conservative
                STREET: Two burner CITY: San Francisco CTATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: linear MOLECULE TYPE: RNA
                                                                                                                                                                                                                                                                                                                                  FILING DATE: 08 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 49
US-10-359-935-41/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CONTINUED DATA:
CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/057,351
FILING DATE: 08-APR-1994
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-CT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/472,802
FILING DATE: 07-4UN-1995
ATTORNEY/AGENT INFORMATION:
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-000821US
TELEPRONE (415) 576-0200
TELEPRAK: (415) 576-0300
TELEFRAK: (415) 576-0300
                                                                                                                                                                                                                                                                                            ADDRESSEE: Townsend and Townsend and Crew LLP
                                                                                                                                                                                                                                                                                                                     STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                Sequence 41, Application US/09057351
Patent No. US20010034439A1
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junii
APPLICANT: Feng, Junii
APPLICANT: Feng, Walter
APPLICANT: Fonk, Walter
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 42, Application US/09057351
Fatent No. US20010034439A1
GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Juni
APPLICANT: Funk, Walter
APPLICANT: Funk, Walter
APLICANT: Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/NS-DOS
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Best Local Similarity 100.
Matches 22; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: RNA
                                                                                                                                                                                                                                                                                                                                                                                                     USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS:
          US-09-057-351-41/c
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US-09-057-351-42/c
                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-057-351-41
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0; Gaps

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FILING DATE: 07-JUN-1995
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LENGTH: 22
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MEDIUM TYPE: Floppy disk
COMPUTER: IS PROPER DEPORT OF THE PROPER DATA:

APPLICATION NUMBER: US/10/359,935
FILING DATE: 07-Feb-2003
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 22;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                               NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 32,944
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFATA: (415) 576-0300
INFORMATION FOR SEQ ID NO: 41:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1;
55;
APPLICATION NUMBER: US/10/359,935
FILING DATE: 07.Feb-2003
CLASSIPICATION: 435
PRIOR APPLICATION: 435
PRIOR APPLICATION NUMBER: US/09/057,351
FILING DATE: 08-APR-1994
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-ULL-1994
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
APPLICATION NUMBER: US 08/472,802
FILING DATE: 07-UN-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 08-APR-1994
PERLICALION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
APPLICATION NUMBER: US 08/330,123
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APPLICATION NUMBER: US/09/057,351
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 27-OCT-1994
APPLICATION NUMBER: US 08/472,802
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear
MOLECULE TYPE: RNA
SEQUENCE DESCRIPTION: SEQ ID NO: 41:
US-10-359-935-41
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US-10-359-935-42/C
US-10-359-935-42/C
Sequence 42, Application US/10359935
Publication No. US20030153076A1
GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
Feng, Junii
Feng, Junii
Feng, Junii
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TYPE: nucleic acid
STRANDEDNESS: single
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STATE: California
COUNTRY: USA
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FULLIANT WEIGHTON:
APPLICANT: Geron Corporation
APPLICANT: Meinrich, Scott
APPLICANT: Meinrich, Scott
APPLICANT: Meinrich, Scott
APPLICANT: Meinrich, Scott
APPLICANT: Lichteteiner, Serge
APPLICANT: Lichteteiner, Serge
APPLICANT: Vaserot, Alain
APPLICANT: Purzan, Ronald
TITLE OF INVENTION: Unhibitors
FILE REPERENCE: 01/006C: 202-12-24
FILE REPERENCE: 01/006C: 202-12-24
FRICK PRILING DATE: 1995-00-4
FRIOR PAPLICATION NUMBER: 08/833,377
PRIOR PAPLICATION NUMBER: 08/833,377
PRIOR PILING DATE: 1999-10-18
FRIOR APPLICATION NUMBER: 09/420,056
FRIOR APPLICATION NUMBER: 09/420,056
FRIOR PILING DATE: 1999-10-18
FRIOR PILING DATE: 1999-10-18
FRIOR PILING DATE: 2000-11-20
NUMBER OF SEQ ID NOS: 11
SEO THE NOS: 11
SEC ID NOS: 11
SEC ID NOS: 11
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100.0%; Pred. No. 55;
ive 0; Mismatches 0; Indele
                                                                                                                                                                                                                                                                                                                                                                                    Length 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Indels
NAME: SCORELLA, JOHN R.
REGISTRATION NUMBER: 32,944
REPERENCE-DOCKET NUMBER: 32,944
REPERENCE TOWNER: 015389-000821US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFATA: (415) 576-0300
INFORMATION FOR SEQ ID NO: 42:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                    DB 1;
55;
                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; Pred. No. 55;
tive 0; Mismatches
                                                                                                                                                                                                                                                                TOPOLOGY: linear MOLECULE TYPE: RNA SEQUENCE DESCRIPTION: SEQ ID NO: 42:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  75
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; Publication No. US20050003404A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 2, Application US/10330872
Publication No. US20030186282A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     54 AACTGAGAAGGGCGTAGGCGCC
                                                                                                                                                                                              LENGTH: 22 base pairs
                                                                                                                                                                                                                    TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 100.0
Matches 22; Conservative
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Best Local Similarity 100.
Matches 22; Conservative
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DB 1; Length 22; 55;
                          RESULT 55
US-10-831-267-13/C
Squence 13, Application US/10831267
Squence 13, Application Wo. US20050009177A1
GENERAL INFORMATION:
APPLICANT: ROWNLEY, Peter T.
TITLE OF INVENTION: TELOWERASE INTERFERENCE
FILE REFERENCE: A-71506-2/RFT/THR
CURRENT APPLICATION NUMBER: US/10/831,267
CURRENT FILING DATE: 2004-04-22
PRIOR FILING DATE: 2002-10-16
PRIOR FILING DATE: 2002-10-22
PRIOR FILING DATE: 2001-10-22
PRIOR PILING DATE: 2001-20-20
PRIOR PILING DATE: 2001-20-20
PRIOR PILING DATE: 2002-03-20
PRIOR PILING DATE: 2002-03-20
PRIOR APPLICATION NUMBER: US 60/359,196
PRIOR APPLICATION NUMBER: US 60/359,196
PRIOR PILING DATE: 2002-03-20
PRIOR PILING DATE: 2002-03-22
NUMBER OF SEQ ID NOS: 23
SOFTWARE: PALENTIN VERSION 3.2
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100.0%; Pred. No. 55,
... 0; Mismatches
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4.9%; Score 22; DB
Best Local Similarity 100.0%; Pred. No. 55;
Matches 22; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19 CTGGGAGGGGTGGTGGCCATTT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Artificial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
Matches 22; Conserva
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ORGANISM: Artificial
                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA
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100.0%; Pred. No. 55;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
4.9%; Score 22; DB 1;
Best Local Similarity 100.0%; Pred. No. 55;
Matches 22; Conservative 0; Mismatches
APPLICANT: Rowley, Peter T.

JITLE OF INVENTION: TELOMERASE INTERFERENCE
FILE REFERENCE: A-71506-1/RFT/THR
CURRENT APPLICATION NUMBER: US/10/831,266
CURRENT FILING DATE: 2004-04-22
RHIOR APPLICATION NUMBER: US/0/33065
PRIOR APPLICATION NUMBER: US 60/345,326
PRIOR APPLICATION NUMBER: US 60/345,326
PRIOR APPLICATION NUMBER: US 60/345,196
PRIOR FILING DATE: 2001-10-22
PRIOR PILING DATE: 2002-02-20
PRIOR PILING DATE: 2002-02-22
PRIOR APPLICATION NUMBER: US 60/383,195
PRIOR PILING DATE: 2002-05-22
NUMBER OF SEQ ID NOS: 17
SOFTWARE: PATENTING DATE: 2002-05-22
NUMBER OF SEQ ID NOS: 17
SOFTWARE: PATENTING DATE: 2002-05-22
NUMBER OF SEQ ID NOS: 17
SOFTWARE: PATENTING DATE: 2002-05-22
NUMBER OF SEQ ID NOS: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 13. Application US/10831266
Publication No. US2005003404A1
GENERAL INFORMATION:
APPLICANT: ROWLEY, PETER T.
TITLE OF INVENTION: TELOMERASE INTERFERENCE
FILE REFERENCE: A-1566-1/RFT/THR
CURRENT APPLICATION NUMBER: US/10/831,266
CURRENT FILING DATE: 2004-04-22
PRIOR PAPLICATION NUMBER: US/10/831,266
PRIOR PAPLICATION NUMBER: US 60/345,326
PRIOR APPLICATION NUMBER: US 60/381,196
PRIOR PILING DATE: 2002-02-20
PRIOR PILING DATE: 2002-05-21
NUMBER OF SEQ 1D NOS: 17
SOFTWARE PRICED NOS: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19 CTGGGAGGGTGGTGGCCATTT 40
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Matches 22; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Artificial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Artificial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 53
US-10-831-266-13/c
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US-10-831-267-12
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                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                            FEATURE
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Length 22; Indels ö

Gaps

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0; Indels

RESULT 56 US-10-811-033-2/c ; Sequence 2, Application US/10811033

; Sequence 12, Application US/10831267; Publication No. US20050009177A1

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CUMUNITY: USB44

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC Compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/10/359,935

FILING DATE: 07-Feb-2003

CLASSIFICATION DATA:

APPLICATION NUMBER: US/09/057,351

FILING DATE: 08-APR-1994

APPLICATION NUMBER: US 08/27,102

FILING DATE: 27-OCT-1994

APPLICATION NUMBER: US 08/472,802

FILING DATE: 27-OCT-1994

APPLICATION NUMBER: US 08/472,802

FILING DATE: 37-OCT-1994

ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4.7%; Score 21; DB 1; Length 21; 00.0%; Pred. No. 64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Indels
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Funk, Walter
Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-000821US
                       APPLICATION NUMBER: US 08/472,802
FILING DATE: 07-07W-1995
ATTORNEY AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REPERENCE/DOCKET NUMBER: 015389-000821US
TELECOMMUNICATION INFORMATION:
TELEFAX: (415) 576-0200
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 25:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Prec. ...
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Publication No. US20030153076A1
GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  184 IGCIGGCCGTTCGCCCCTCC 204
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21 TGCTGGCCCGTTCGCCCCTCC 1
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COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 100.1
Matches 21; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS
LENGTH: 21 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: nucleic acid
STRANDEDNESS: singl
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-10-359-935-25/c
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                                                  APPLICANT: Geron Corporation
APPLICANT: Meinrich, Scott
APPLICANT: Atkinson III, Edward
APPLICANT: Atkinson III, Edward
APPLICANT: Atkinson III, Edward
APPLICANT: Lichtsteiner, Serge
APPLICANT: Drivan, Ronald
TITLE OF INVENTION: Using Purified Telomerase to Identify Telomerase Activators and
TITLE OF INVENTION: Using Purified Telomerase to Identify Telomerase Activators
FILE REFERENCE: 011/006C
CURRENT FILING DATE: 2004-03-26
PRIOR APPLICATION NUMBER: 08/10/330,872A
PRIOR PLING DATE: 1995-08-04
PRIOR PLING DATE: 1997-04-04
PRIOR PLING DATE: 1997-04-04
PRIOR PLING DATE: 1999-10-18
PRIOR FILING DATE: 1997-04-04
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COUNTRY: USA
ZIP: 9411-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/057,351
TT.ING DATE: 08-ARR-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             E: Townsend and Townsend and Crew LLP Two Embarcadero Center, Eighth Floor
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CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CTAACCCTAACTGAGAAGGGCG 67
Publication No. US20050089883A1
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STATE: California
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ORGANISM: Homo sapiens
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Publication No. US20050153916A1
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                                                                                                      Gaps
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                                                                            Query Match
4.7%; Score 21; DB 1; Length 21;
Best Local Similarity 100.0%; Pred. No. 64;
Matches 21; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
4.7%; Score 21; DB 1; Length 21;
Best Local Similarity 81.0%; Pred. No. 64;
Matches 17; Conservative 4; Mismatches 0; Indels
          TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: DNA
SEQUENCE DESCRIPTION: SEQ ID NO: 25:
                                                                                                                   184 TGCTGGCCCGTTCGCCCTCC 204
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 21 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: RNA
ORGANISM: Artificial Sequence
                                                          US-10-359-935-25
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US-10-923-330-537
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US-10-923-330-542/c
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Sequence 539, Application US/10923330

Sequence 539, Application US/10923330

GENERAL INFORMATION:

APPLICANT: Weskiggen, James

APPLICANT: Weskiggen, Leonid

TITLE OF INWENTION: Expression Using Short Interfering RNA (siNA)

FILE REFERENCE: 400/209 (MBHE02-708-C)

CURRENT APPLICATION NUMBER: US/10/923,330

CURRENT FILING DATE: 2004-08-20

FRIOR PELING DATE: 2004-08-20

PRIOR FILING DATE: 2004-06-24

PRIOR FILING DATE: 2004-05-24

PRIOR FILING DATE: 2004-05-24

PRIOR APPLICATION NUMBER: US 10/750,803

PRIOR FILING DATE: 2003-11-24

PRIOR APPLICATION NUMBER: US 10/757,803

PRIOR APPLICATION NUMBER: US 10/753,803

PRIOR APPLICATION NUMBER: US 10/753,803
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, OTHER INFORMATION: Description of Artificial Sequence: sinA sense region
US-10-923-330-539
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: Description of Artificial Sequence: siNA sense region
PRIOR APPLICATION NUMBER: US 10/720,448

PRIOR FILING DATE: 2003-11-24

PRIOR PILING DATE: 2003-11-23

PRIOR PILING DATE: 2003-11-23

PRIOR APPLICATION NUMBER: US 10/444,853

PRIOR APPLICATION NUMBER: US 10/444,853

PRIOR APPLICATION NUMBER: PCT/US03/05346

PRIOR PILING DATE: 2003-02-20

PRIOR PILING DATE: 2003-02-20

PRIOR PILING DATE: 2003-02-0

PRIOR PILING DATE: 2003-02-0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1; Mismatches
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ORGANISM: Artificial Sequence
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Matches 20; Conserva
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LENGTH: 21
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; OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region US-10-923-330-541
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Publication No. US20050153916A1
GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Beigelman, Jemes
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: RAY Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Expression Using Short Interfering RNA (81NA)
                           Gaps
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Score 21; DB 1; Length 21;
Pred. No. 64;
4; Mismatches 0; Indels
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64;
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                                                                                                                                       Sequence 541, Application US/10923330 Publication No. US20050153916A1 GENERAL INFORMATION:
                                                   397 GCGCGCGATTCCCTGAGCTG 417
                                                                   1 GCGGCGCAUUCCCUGAGCUG 21
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  Query Match
Best Local Similarity 81.0%;
Matches 17; Conservative
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Gaps

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RESULT 66 SEPTIONE 544/C

SEQUENCE 546/C

SEQU
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         PRIOR FILING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR FILING DATE: 2003-02-20
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 768
SOFTWARE: Patentin version 3.3
LENGTH: 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              283 GCACCCACTGCCACCGCGAAG 303
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                                                                                                                                                                                                                                                                                                                                                               TYPE: RNA
ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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Matches 21; Conserva
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Sequence 543, Application US/10923330

Publication No. US20050153916A1

GENERAL INFORMATION:

APPLICANT: Sirna Therapeutics, Inc.

APPLICANT: McSwiggen, James

APPLICANT: Beigelman, Leonid

TILLE OF INVENTION: RNA INterference Mediated Inhibition of Telomerase Gene

TILLE OF INVENTION: Expression Using Short Interfering RNA (siNA)

FILE REFERENCE: 400/209 (MBHB02-708-C)

CURRENT PILING DATE: 2004-08-20

PRIOR APPLICATION NUMBER: PCT/US03/04088

PRIOR APPLICATION NUMBER: PCT/US03/04088

PRIOR APPLICATION NUMBER: DCT/US04/16390

PRIOR FILING DATE: 2004-05-24

PRIOR APPLICATION NUMBER: US 10/757,803

PRIOR FILING DATE: 2004-04-16

PRIOR FILING DATE: 2004-04-16

PRIOR FILING DATE: 2004-01-12

PRIOR FILING DATE: 2003-01-124

PRIOR FILING DATE: 2003-01-123

PRIOR FILING DATE: 2003-01-123

PRIOR FILING DATE: 2003-01-23

PRIOR FILING DATE: 2003-01-23

PRIOR FILING DATE: 2003-05-23

PRIOR PRIOR APPLICATION NUMBER: US 10/444,853

PRIOR FILING DATE: 2003-05-23

PRIOR FILING DATE: 2003-05-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; OTHER INFORMATION: Description of Artificial Sequence: siNA antisense region US-10-923-330-542
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             CURRENT APPLICATION NUMBER: US10/923,330

CURRENT APPLICATION NUMBER: PC1/US03/04088

PRIOR APPLICATION NUMBER: PC7/US03/04088

PRIOR FILING DATE: 2004-05-24

PRIOR FILING DATE: 2004-05-24

PRIOR FILING DATE: 2004-05-24

PRIOR FILING DATE: 2004-05-24

PRIOR PLILING DATE: 2004-01-14

PRIOR FILING DATE: 2004-04-16

PRIOR FILING DATE: 2004-04-16

PRIOR PLING DATE: 2003-11-24

PRIOR PLING DATE: 2003-11-24

PRIOR PLING DATE: 2003-11-23

PRIOR PLING DATE: 2003-11-23

PRIOR PLING DATE: 2003-05-23

PRIOR PLING DATE: 2003-05-20

PRIOR PLING DATE: 2003-05-30

PRIOR PLING DATE: 20
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4.7%; Score 21; DB 1; Length 21;
Best Local Similarity 100.0%; Pred. No. 64;
Matches 21; Conservative 0; Mismatches 0; Indels
(MBHB02-708-C)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: RNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 65
US-10-923-330-543/c
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COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              41 TTTGTCTAACCCTAACTGAG 60
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               , MOLECULE TYPE: RNA
US-09-057-351-40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-10-359-935-7/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: USA
ZIP: 9411-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC Compatible
OPERATING SYSTEM: FC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/057,351
FILING DATE: 08-APR-1994
FILING DATE: 07-JUL-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 07-JUL-1994
PRIOR APPLICATION NUMBER: US 08/330,123
FILING DATE: 2-OCT-1994
PRIOR APPLICATION NUMBER: 32,944
FRIENT APPLICATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 32,944
REDEPAN: (415) 5.76-0300
RELEPAN: (415) 5.76-0300
RELEPAN: MARCHERICE/REVERISTICS: SEQUENCE CHARACTERISTICS: SEQUENCE CHARACTERISTICS: SEGUENCE CHARACTERISTIC
                                                                                                                                       Sequence 7, Application US/09057351
; Sequence 7, Application US/09057351
; Patent No. US20010034439A1
; GENERAL INFORMATION:
APPLICANT: Funk, Walter
APPLICANT: Funk, Walter
APPLICANT: Funk, Walter
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
ADDRESSE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
COUNTRY: California
COUNTRY: USA
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Feng, Junli
Funk, Walter
Andrews, William H.
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Patent No. US200100134439A1
GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junit
APPLICANT: Feng, Walter
APPLICANT: Funk, Walter
APPLICANT: APAGREWS, William H.
21 GCGCGCGCGATTCCCTGAGC 1
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STRANDEDNESS: sing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
Matches 20; Conserva
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US-09-057-351-40/c
                                                                                                                                    US-09-057-351-7/c
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Gaps
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                                                                                                                                                                               COUNTRY: USA

ZIP: 9411-3834

COMPUER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATION SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTION DATA:
APPLICATION NDATA:
APPLICATION NDATA:
APPLICATION NDATA:
APPLICATION NDATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 09-APR-1994
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: O'-JUL-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/472,802
FILING DATE: O'-JUL-1994
APPLICATION NUMBER: US 08/472,802
FILING DATE: O'-JUN-1995
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 015389-000821US
TELECOMMUNICATION INFORMATION:
TELEBROME: (415) 576-0200
TELEBROME: (415) 576-0300
INFORMATION FOR SEQ ID NO: 40:
SEQUENCE CHARACTERISTICS:
LENGTH: C. D DASE DATES
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Feng, Junli
Funk, Walter
Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
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APPLICATION NUMBER: US 08/330,123
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100.0%; Prec
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                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO: 40: SEQUENCE CHARACTERISTICS: LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            41 TITGICTAACCCTAACTGAG 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20 TTTGTCTAACCCTAACTGAG 1
                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              76 GIGCITITGCICCCGCGC
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Best Local Similarity 100.0
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 100.
Matches 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-09-018-125-2/c
                                                                                                                                                                                                                                                                                                                                                                                                                 US-10-359-935-40
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        CURRENT APPLICATION DATA:

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/10/359,935

FILING DATE: 07-Feb-2003

PRIOR APPLICATION NUMBER: US/09/057,351

FILING DATE: 04-APR-1994

APPLICATION NUMBER: US 08/272,102

FILING DATE: 07-JUL-1994

APPLICATION NUMBER: US 08/30,123

FILING DATE: 07-JUL-1994

APPLICATION NUMBER: US 08/30,123

FILING DATE: 07-JUL-1995

ATORNEY/AGENT INFORMATION:

NAME: STORELLA JOHN R.

RESISTRATION NUMBER: 32,944

REFERENCE/DOCKET NUMBER: 32,944

REFERENCE/DOCKET NUMBER: 32,944

REFERENCE/DOCKET NUMBER: 1576.0200

TELEEPHONE: (415) 576.0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4.4%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 73; 0; Indels tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco STATE: California COUNTRY: USA ZIP: 94111-13834 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk COMPUTER: IBM PC COMPUTER: IBM PC COMPUTER: DO PERATING SYSTEM: PC-DOS/MS_DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/359,935 FILING DATE: 07-Feb-2003 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Feng, Junli
Funk, Walter
Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/057,351
FILING DATE: 08-APP-1994
APPLICATION NUMBER: US 08/272,102
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; TOPOLOGY: Linear; MOLECTLE TYPE: DNA SEQUENCE DESCRIPTION: SEQ ID NO: 7: US-10-359-935-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-10-359-935-40/c
US-10-359-935-40/c
; Sequence 40, Application US/10359935
; Publication No. US20030153076A1
; GENERAL INFORMATION:
; APPLICANT: Villeponteau, Bryant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 07-JUL-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                   (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GGTTGCGGAGGGTGGGCCTG 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20 GGTTGCGGAGGGTGGGCCTG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20; Conservative
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Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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GENERAL INFORMATION:

APPLICANT: Silverman, Robert H.

APPLICANT: Silverman, Robert H.

APPLICANT: Cowell, John K.

APPLICANT: Cowell, John K.

APPLICANT: Cowell, John K.

APPLICANT: Torrence, Paul F.

TITLE OF INVENTION: RNASE L ACTIVATORS AND ANTISENSE OLIGONUCLEOTIDES

TITLE OF INVENTION: RNASE L (18/09/018,125A)

TITLE OF INVENTION: BFFECTIVE TO TREAT TELOMERASE-EXPRESSING MALIGNANCIES

FILE REFERENCE: 8656-022

CURRENT APPLICATION NUMBER: (08/09/018,125A)

CURRENT FILING DATE: 1999-02-03

EARLIER APPLICATION NUMBER: 60/044,507

EARLIER FILING DATE: 1997-04-21

NUMBER OF SEQ ID NOS: 9

SEQ ID NO 2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Indels
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Pred. No. 82;
0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ) OTHER INFORMATION: Description of Artificial Sequence:
) OTHER INFORMATION: oligonucleotide
US-09-018-125-2
                                                                 ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-000821US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4.4%; Score 20; DB 100.0%; Pred. No. 73; Live 0; Mismatches
FILING DATE: 27-OCT-1994
APPLICATION NUMBER: US 08/472,802
FILING DATE: 07-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear
MOLECULE TYPE: RNA
SEQUENCE DESCRIPTION: SEQ ID NO: 40:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 72
US-10-016-490C-8
; Sequence 8, Application US/10016490C
; GENERAL INFORMATION:
; GENERAL INFORMATION:
; APPLICANT: Yin, James Q.
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ORGANISM: Artificial
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                                                                                                                                                                                                                                                                                                                          US-10-831-267-17
                                                                                                                                                                                                                                TYPE: RNA
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TITLE OF INVENTION: Methods for design and selection of short double-stranded TITLE OF INVENTION: oligonucleotides, and compounds of gene drugs PILE REFERENCE: 01-27-91 CURRENT APPLICATION NUMBER: US/10/016,490C CURRENT FILING DATE: 2002-11-22 NUMBER OF SEQ ID NOS: 51 SOFTWARE: Patentin version 3.1 SEQ ID NO 8
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                                                                                                                                                                                                                                                                                                                                                Length 19;
                                                                                                                                                                                                                                                                                                                                                                                          0; Indels
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4.2%; Score 19; DB 1;
Best Local Similarity 100.0%; Pred. No. 82;
Matches 19; Conservative 0; Mismatches
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Sequence 16, Application US/10831267

Sequence 16, Application US/10831267

Publication No. US20050009177A1

GENERAL INFORMATION:
TITLE OF INVENTION: TELOMERASE INTERFERENCE
FILE REPRENCE: A-71506-2/RRT/THR

CURRENT APPLICATION NUMBER: US/10/831,267

CURRENT APPLICATION NUMBER: US/10/831,267

CURRENT APPLICATION NUMBER: US 60/345,326

PRIOR PILING DATE: 2004-04-22

PRIOR PILING DATE: 2004-16

PRIOR FILING DATE: 2001-16

PRIOR FILING DATE: 2001-16

PRIOR APPLICATION NUMBER: US 60/359,196

PRIOR PILING DATE: 2001-10-22

PRIOR APPLICATION NUMBER: US 60/359,196

PRIOR FILING DATE: 2002-02-20

NUMBER OF SEQ ID NOS: 23

SOFTWARE: PARCH NOS: 23

SEQ ID NO 16

LENGTH: 19
                                                                                                                                                                                                                                                                           CTHER INFORMATION: The same as those in human US-10-016-490C-8
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; Publication No. US20050009177A1
; GENERAL INFORMATION:
; APPLICATY: ROWLEY, Peter T.
; TILE OF INVENTION: TELOMERASE INTERFERENCE
; FILE REFERENCE: A-71506-2/RFT/THR
; CURRENT APPLICATION NUMBER: US/10/831,267
; CURRENT FILING DATE: 2004-04-22
; PRIOR APPLICATION NUMBER: PCT/US 02/33146
; PRIOR PILING DATE: 2002-10-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                          372 AGAGGAACGGAGCGAGTCC 390
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 AGAGGAACGGAGCGAGTCC 19
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                                                                                                                                                                                                                                ORGANISM: Artificial Sequence
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Artificial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 74
US-10-831-267-17/c
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                                                                                                                                                                                                             TYPE: DNA
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82;
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82;
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; Sequence 18, Application US/10831267
; Publication No. US20050009177A1
; GENERAL INFORMATION:
    APPLICANT: Rowley, Peter T.
    TITLE OF INVENTION: TELOMERASE INTERFERENCE
; TITLE OF INVENTION: TELOMERASE INTERFERENCE
; CURRENT ELING DATE: 2004-04-22
    PRIOR PILING DATE: 2004-04-22
; PRIOR APPLICATION NUMBER: PCT/US 02/33146
; PRIOR APPLICATION NUMBER: DCT/US 02/33146
; PRIOR PLING DATE: 2001-10-22
; PRIOR PLING DATE: 2001-10-22
; PRIOR FILING DATE: 2002-20
; PRIOR FILING DATE: 2002-05-20
; PRIOR FILING DATE: 2002-05-20
; PRIOR PILING DATE: 2002-05-20
; PRIOR PILING DATE: 2002-05-20
; RIOR APPLICATION NUMBER: US 60/383,195
; SOFURMARE: Patentin Version 3.2
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                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
4.2%; Score 19; DB
Best Local Similarity 100.0%; Pred. No. 82;
Matches 19; Conservative 0; Mismatches
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4.2%; Score 19; DB
Best Local Similarity 100.0%; Pred. No. 82;
Matches 19; Conservative 0; Mismatches
PRIOR APPLICATION NUMBER: US 60/345,326
PRIOR FILING DATE: 2001-10-22
PRIOR APPLICATION NUMBER: US 60/359,196
PRIOR FILING DATE: 2002-02-20
PRIOR FILING DATE: 2002-05-22
NUMBER OF SEQ ID NOS: 23
SOFTWARE: Patentin version 3.2
SEQ ID NO 17
LENGTH: 19
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FEATURE:
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82;
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82;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: BARK: Joffre
APPLICANT: BARK: Joffre
APPLICANT: Gronin, Maureen
APPLICANT: BARK: Joffre
APPLICANT: BARK: Jose
TITLE OF INVENTION: GENE EXPRESSION PROFILING OF EGFR
TITLE OF INVENTION: POSITIVE CANCER
FILE REFERENCE: 39740-0005
CURRENT APPLICATION NUMBER: US/10/714,195
CURRENT PILING DATE: 2003-11-14
PRIOR APPLICATION NUMBER: 60/427090
PRIOR FILING DATE: 2003-11-15
NUMBER OF SEQ ID NOS: 372
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 343
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NS-10-714-195-344/c

Sequence 344, Application US/10714195

Publication No. US2005019785A1

GENERAL INFORMATION:

APPLICANT: Baker, Joffre

APPLICANT: Croin, Maureen

APPLICANT: Baselga, Jose

TILLE OF INVENTION: GENE EXPRESSION PROFILING OF EGFR

TILLE OF INVENTION: POSITIVE CANCER

FILE REFERENCE: 39740-0005
                                                                                                                                                                                                                                                                                                                                                       Query Match
4.2%; Score 19; DB
Best Local Similarity 100.0%; Pred. No. 82;
Matches 19; Conservative 0; Mismatches
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4.2%; Score 19; DB
Best Local Similarity 100.0%; Pred. No. 82;
Matches 19; Conservative 0; Mismatches
               PRIOR APPLICATION NUMBER: US 60/345,326
PRIOR FILING DATE: 2001-10-22
PRIOR APPLICATION NUMBER: US 60/359,196
PRIOR APPLICATION NUMBER: US 60/383,195
PRIOR FILING DATE: 2002-02-20
NUMBER OF SEQ ID NOS: 23
SOFTWARE: PatentIn version 3.2
SEQ ID NO 19
LENGTH: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 343, Application US/10714195
Publication No. US20050019785A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                          OTHER INFORMATION: oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                       270 GGCTTCTCCGGAGGCACCC 288
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PRIOR FILING DATE: 2002-10-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; OTHER INFORMATION: primer US-10-714-195-343
                                                                                                                                                                                                                             TYPE: DNA
ORGANISM: Artificial
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; OTHER INFORMATION: Description of Artificial Sequence: Target Sequence/sinA sense re
US-10-923-330-7
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Publication No. US20050153946A1

GENERAL INFORMATION:

APPLICANT: Sirna Therapeutics, Inc.

APPLICANT: Baigadiann, Leonid

TITLE OF INVENTION: RAD. Thereference Mediated Inhibition of Telomerase Gene

TITLE OF INVENTION: RAD. Thereference Mediated Inhibition of Telomerase Gene

TITLE OF INVENTION: RAD. Thereference Mediated Inhibition of Telomerase Gene

TITLE OF INVENTION: Baptession Using Short Interfering RNA (sinA)

FILE REFERENCE: 400.0209 (MEMBERS: US/10/923,330

CURRENT FILING DATE: 2004-08-20

FRIOR APPLICATION NUMBER: US 60/396,600

FRIOR APPLICATION NUMBER: US 10/926,966

FRIOR PELING DATE: 2004-04-16

FRIOR PELING DATE: 2004-04-16

FRIOR FILING DATE: 2004-01-14

FRIOR FILING DATE: 2004-01-14

FRIOR FILING DATE: 2003-11-24

FRIOR FILING DATE: 2003-11-24

FRIOR FILING DATE: 2003-11-23

FRIOR APPLICATION NUMBER: US 10/750,803

FRIOR FILING DATE: 2003-11-23

FRIOR APPLICATION NUMBER: US 10/750,803

FRIOR PELING DATE: 2003-11-23

FRIOR PELING DATE: 2003-11-23

FRIOR APPLICATION NUMBER: PCT/US03/05028

FRIOR FILING DATE: 2003-02-20

FRIOR FILING DATE: 2003-02-20
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NUMBER OF SEQ ID NOS: 768
SOFTWARE: PatentIn version 3.3
SEQ ID NO 7
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4.2%; Score 19; DB 1; Length 19;
Best Local Similarity 100.0%; Pred. No. 82;
Matches 19; Conservative 0; Mismatches 0; Indels
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4.2%; Score 19; DB 1; Length 19;
Best Local Similarity 78.9%; Pred. No. 82;
Matches 15; Conservative 4; Mismatches 0; Indels
CURRENT FILING DATE: 2003-11-14
PRIOR APPLICATION NUMBER: 60/427090
PRIOR FILING DATE: 2003-11-15
NUMBER OF SEQ ID NOS: 372
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 344
LENGTH: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           431 CAGGACTCGGCTCACACAT 449
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                                                                                                                                                                                                                                                                                          TYPE: DNA
ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                          ; OTHER INFORMATION: primer US-10-714-195-344
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OTHER INFORMATION: Description of Artificial Sequence: Target Sequence/#1NA sense re
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APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Bodgelman, James
APPLICANT: Baggelman, James
APPLICANT: Baggelman, Jeonid
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE REPERBENCE: 400/209 (WHBDED-708-C)
CURRENT APPLICATION NUMBER: US 10/4023, 330
CURRENT FILING DATE: 2004-05-24
PRIOR APPLICATION NUMBER: US 60/396, 600
PRIOR FILING DATE: 2004-05-24
PRIOR FILING DATE: 2004-04-16
PRIOR FILING DATE: 2004-04-16
PRIOR APPLICATION NUMBER: US 10/426, 966
PRIOR FILING DATE: 2004-01-14
PRIOR APPLICATION NUMBER: US 10/693, 059
PRIOR FILING DATE: 2003-11-24
PRIOR FILING DATE: 2003-11-24
PRIOR PAPLICATION NUMBER: US 10/693, 059
PRIOR FILING DATE: 2003-11-23
PRIOR FILING DATE: 2003-11-23
PRIOR PAPLICATION NUMBER: US 10/693, 059
PRIOR FILING DATE: 2003-11-23
PRIOR PAPLICATION NUMBER: PCT/US03/05346
PRIOR FILING DATE: 2003-02-20
PRIOR PLING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR PLING DATE: 2003-02-20
PRIOR PRIOR PLING DATE: 2003-02-20
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PCT/US04/16390
                                      FILING DATE: 2004-05-24
APPLICATION NUMBER: US 10/826,966
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 10, Application US/10923330 Publication No. US20050153916A1 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: RNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity 84.2
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-10-923-330-9
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US-10-923-330-8
                                                                                                                                                          US-10-922-330-8

US-10-922-330-8

Sequence 8, Application US/10923330

Publication No. US20050153916A1

GENERAL INPORATION:

APPLICANT: SITRA Therapeutics, Inc.

APPLICANT: McSw4gen, James

APPLICANT: WNWBER: US/10/923,330

CURRENT APPLICATION NUMBER: US/10/923,330

FRIOR FILING DATE: 2004-05-24

FRIOR FILING DATE: 2004-05-24

FRIOR FILING DATE: 2004-05-24

FRIOR FILING DATE: 2004-01-14

FRIOR PLICATION NUMBER: US/10/59,666

FRIOR APPLICATION NUMBER: US/10/59,666

FRIOR FILING DATE: 2003-11-23

FRIOR FILING DATE: 2003-12-20

FRIOR PELICATION NUMBER: PCT/US03/05028

FRIOR PELICATION NUMBER: PCT/US03/05028

FRIOR PELICATION NUMBER: PCT/US03/05028

FRIOR FILING DATE: 2003-02-20

FRIOR FILING DATE: 2003-02-20

FRIOR FILING DATE: 2003-02-20

FRIOR FILING DATE: 2003-02-20

FRIOR FILING DATE: 2003-02-30

FRIOR FILING DATE: 2003-03-31

FRIOR FILING DATE: 2003-03-33

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Sequence 9, Application US/10923330;
Publication No US20050153916A1;
GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Biggen, James
APPLICANT: Beigen, James
APPLICANTON: Expression Using Short Interfering RNA (sinA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
FILE REFERENCE: 400/209 (MBHB02-708-C)
CURRENT APPLICATION NUMBER: DCT/US03/04088
PRIOR FILING DATE: 2004-08-20
PRIOR FILING DATE: 2004-05-24
PRIOR FILING DATE: 2002-07-17
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   1 UGGGCCUGGGAGGGUGGU 19
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ORGANISM: Artificial Sequence
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OTHER INFORMATION: Description of Artificial Sequence: Target Sequence/sinA sense re
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, James
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RAPICASION WHEBER: US/10/923,330
CURRENT APPLICATION WHEBER: US/10/923,330
CURRENT APPLICATION WHEBER: US/20/96-20
PRIOR FILING DATE: 2004-06-24
PRIOR PLING DATE: 2004-05-24
PRIOR APPLICATION WHERE: US 10/757,803
PRIOR APPLICATION WHERE: US 10/757,803
PRIOR APPLICATION WHERE: US 10/757,803
PRIOR PLING DATE: 2004-01-14
PRIOR PLING DATE: 2003-01-24
PRIOR PLING DATE: 2003-01-24
PRIOR PLING DATE: 2003-11-23
PRIOR PLING DATE: 2003-11-23
PRIOR PLING DATE: 2003-11-23
PRIOR PLING DATE: 2003-11-23
PRIOR PLING DATE: 2003-01-24
PRIOR PLING DATE: 2003-01-24
PRIOR PLING DATE: 2003-01-24
PRIOR PLING DATE: 2003-01-23
PRIOR PLING DATE: 2003-02-20
PRIOR PLING 
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Sequence 13, Application US/10923330

Sequence 13, Application US/10923330

PUBLIcature No. US20050153916A1

GENERAL INFORMATION:

APPLICANT: Sirna Therapeutics, Inc.

APPLICANT: Beigalman, Leonid

TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)

TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)

TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)

TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)

CURRENT FILING DATE: 2004-08-20

PRIOR PALICATION NUMBER: US 60/396, 600

PRIOR PAPLICATION NUMBER: US 60/396, 600

PRIOR APPLICATION NUMBER: US 60/396, 600

PRIOR PAPLICATION NUMBER: US 10/826,966

PRIOR FILING DATE: 2004-05-24

PRIOR FILING DATE: 2004-04-16

PRIOR PILING DATE: 2004-04-16

PRIOR PILING DATE: 2004-01-14

PRIOR PILING DATE: 2004-01-14

PRIOR PILING DATE: 2004-01-14
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4.2%; Score 19; DB 1; Length 19;
Best Local Similarity 73.7%; Pred. No. 82;
Matches 14; Conservative 5; Mismatches 0; Indels
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ORGANISM: Artificial Sequence
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                                                                           ; OTHER INFORMATION: Description of Artificial Sequence: Target Sequence/siNA sense
US-10-923-330-10
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US-10-923-330-11
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4.2%; Score 19; DB 1; Length 19,
Best Local Similarity 63.2%; Pred. No. 82;
Matches 12; Conservative 7; Mismatches 0; Indels
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4.2%; Score 19; DB 1; Length 19;
Best Local Similarity 63.2%; Pred. No. 82;
Matches 12; Conservative 7; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                         68 TAGGCGCCGTGCTTTTGCT 86
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TYPE: RNA
ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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US-10-923-330-12
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                                                    FEATURE:
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Target Sequence/siNA sense r
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APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Bisna Therapeutics, Inc.
APPLICANT: Beigelman, Leonid
PUTLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT APPLICATION NUMBER: PCT/US3/04088
PRIOR FILING DATE: 2004-05-24
PRIOR FILING DATE: 2004-05-24
PRIOR FILING DATE: 2004-05-24
PRIOR FILING DATE: 2004-05-24
PRIOR FILING DATE: 2004-01-14
PRIOR FILING DATE: 2004-01-14
PRIOR FILING DATE: 2004-01-14
PRIOR FILING DATE: 2004-01-14
PRIOR FILING DATE: 2003-01-12
PRIOR FILING DATE: 2003-01-12
PRIOR FILING DATE: 2003-01-12
PRIOR PRIOR PRIOR DATE: 2003-01-13
PRIOR FILING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: US 10/693,059
PRIOR PRILING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR PLING DATE: 2003-02-20
PRIOR PRILING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR PLING DATE: 2003-02-20
PRIOR PRILING DATE: 2003-02-20
PRIOR PRILING DATE: 2003-02-20
PRIOR PRILING DATE: 2003-02-20
PRIOR PRILING DATE: 2003-03-3-3
PRIOR PLING DATE: 2003-02-20
PRIOR PRILING DATE: 2003-03-3-3
PRIOR PRILING DATE: 2003-05-3-3

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Publication No. US20050153916A1
GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Beigelman, Leonid
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: EXPRESSION USING Short Interfering RNA (SINA)
FILE REFERENCE: 400/209 (MBHB02-708-C)
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         Length 19;
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84.2%; Pred. No. 82;
tive 3; Mismatches 0; Indels
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US-10-923-330-15
         Score 19; DB 1;
Pred. No. 82;
                                                                                                        5; Mismatches
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Sequence 15, Application US/10923330
Publication No. US20050153916A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                             1 GCCGCCUUCCACCGUUCAU 19
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ORGANISM: Artificial Seguence
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Best Local Similarity 84.2'
Matches 16; Conservative
                                                                                                             Conservative
         Query Match
Best Local Similarity
Matches 14; Conserv
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US-10-923-330-14
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TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE REFERENCE: 400/209 (NBHB02-708-C)
CURRENT APPLICATION NUMBER: 2004-08-20
PRIOR PILING DATE: 2004-08-20
PRIOR FILING DATE: 2004-05-24
PRIOR FILING DATE: 2004-05-24
PRIOR PLING DATE: 2004-05-24
PRIOR FILING DATE: 2004-05-24
PRIOR FILING DATE: 2004-05-24
PRIOR FILING DATE: 2004-06-14
PRIOR FILING DATE: 2004-01-14
PRIOR FILING DATE: 2004-01-14
PRIOR FILING DATE: 2003-11-24
PRIOR FILING DATE: 2003-05-23
PRIOR FILING DATE: 2003-05-23
PRIOR FILING DATE: 2003-02-20
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PRIOR FILING DATE: 2003-11-24
PRIOR PELICATION NUMBER: US 10/693,059
PRIOR FILING DATE: 2003-11-23
PRIOR PELICATION NUMBER: US 10/444,853
PRIOR PELICATION NUMBER: PCT/US03/05346
PRIOR PILING DATE: 2003-05-23
PRIOR PILING DATE: 2003-02-20
PRIOR PILING DATE: 2003-02-30
PRIOR PRIOR PRIOR APPLICATION DATE: 2003-02-30
PRIOR PILING DATE: 2003-02-20
PRIOR PILING DATE: 20
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Publication No. US20050153916A1
GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
APPLICANT: Reigelman, Leonid
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ORGANISM: Artificial Sequence
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Matches 17; Conservative
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OTHER INFORMATION: Description of Artificial Sequence: Target Sequence/sink sense re
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US-10-923-330-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 18, Application US/10923330

Publication No. US20050153916A1

GRENEAL INFORMATION:

APPLICANT: Sirna Therapeutics, Inc.

APPLICANT: McSwiggen, James

APPLICANT: Beigelman, Leonid

TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)

TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)

FILE REPERENCE: 400/209 (MBHB02-708-C)

CURRENT APPLICATION NUMBER: US/10/923,330

CURRENT APPLICATION NUMBER: US/00/923,330

GURENT APPLICATION NUMBER: PCT/US03/04088

PRIOR FILING DATE: 2004-06-20

PRIOR FILING DATE: 2004-05-40

PRIOR PILING DATE: 2004-05-40

PRIOR PLING DATE: 2004-01-14

PRIOR PLING DATE: 2004-01-14

PRIOR PLING DATE: 2004-01-14

PRIOR PLING DATE: 2004-01-14

PRIOR PLING DATE: 2003-11-23

PRIOR APPLICATION NUMBER: US 10/757,803

PRIOR APPLICATION NUMBER: US 10/757,803

PRIOR PLING DATE: 2003-11-23

PRIOR PLING DATE: 2003-11-23

PRIOR PLING DATE: 2003-05-23

PRIOR PLING DATE: 2003-05-23
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             PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR FILING DATE: 2003-02-20
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 768
SOFTWARE: Patentin version 3.3
SEQ ID NO 17
LENGTH: 19
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SOFTWARE: PatentIn version 3.3
SEQ ID NO 18
LENGTH: 19
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Pred. No. 82;
3; Mismatches 0; Indels
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                                                                                                                                                                                               TYPE: RNA
ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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Best Local Similarity 84.2%;
Matches 16; Conservative
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Publication No. US20050153916A1
GENERAL INFORMATION:
Hubblication No. US20050153916A1
GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
CURRENT APPLICATION: Expression Using Short Interfering RNA (siNA)
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT FILING DATE: 2004-06-20
PRIOR FILING DATE: 2004-05-24
PRIOR APPLICATION NUMBER: US 10/826,966
PRIOR FILING DATE: 2004-06-14
PRIOR FILING DATE: 2004-01-14
PRIOR FILING DATE: 2004-01-14
PRIOR FILING DATE: 2003-11-24
PRIOR FILING DATE: 2003-11-24
PRIOR FILING DATE: 2003-11-24
PRIOR FILING DATE: 2003-11-24
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PRIOR FILING DATE: 2003-05-23
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CURRENT APPLICATION NUMBER: US/10/923,330
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APPLICATION NUMBER: PCT/US03/05346
FILING DATE: 2003-02-20
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1 AUGUCAGCUGCUGGCCCGU 19
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ORGANISM: Artificial Sequence
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Matches 14; Conserv
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; OTHER INFORMATION: Description of Artificial Sequence: Target Sequence/sinA sense re
US-10-923-330-20
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UNERRY APPLICATION NUMBER: US/10/923,330

UURRENT FILING DATE: 2004-08-20

UURRENT PILING DATE: 2004-08-20

UURRENT PILING DATE: 2004-08-20

UURRENT PILING DATE: 2004-08-24

PRIOR PILING DATE: 2004-08-24

PRIOR PILING DATE: 2004-08-24

PRIOR PILING DATE: 2004-09-17

PRIOR PILING DATE: 2004-04-16

PRIOR PILING DATE: 2004-01-14

PRIOR PILING DATE: 2004-01-14

PRIOR PILING DATE: 2004-01-14

PRIOR PILING DATE: 2004-01-14

PRIOR PILING DATE: 2003-11-23

PRIOR PILING DATE: 2003-02-20

PRIOR PI
                                 FRIOR FILING DATE: 2004-04-15
PRIOR FILING DATE: 2004-04-15
PRIOR PELICATION NUMBER: US 10/757,803
PRIOR PELICATION NUMBER: US 10/757,803
PRIOR FILING DATE: 2003-01-124
PRIOR FILING DATE: 2003-11-24
PRIOR FILING DATE: 2003-11-24
PRIOR FILING DATE: 2003-10-30-3
PRIOR FILING DATE: 2003-05-33
PRIOR FILING DATE: 2003-05-32
PRIOR FILING DATE: 2003-05-32
PRIOR PILING DATE: 2003-02-30
PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR PILING DATE: 2003-02-30
PRIOR PILING DATE: 2003-02-30
PRIOR PILING DATE: 2003-02-30
PRIOR FILING DATE: 2003-02-30
PRIOR FILING DATE: 2003-02-30
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PRIOR PILING DATE: 2003-02-30
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US 10/826,966
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SEQ ID NO 21
LENGTH: 19
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US-10-923-330-19
                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
APPLICANT: Beigglaman, Leonia
TITLE OF INVENTION: RAY Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE REFERENCE: 400/209 (MBHB02-708-C)
CURRENT APPLICATION NUMBER: US/10/323,330
CURRENT APPLICATION NUMBER: US 60/396,600
PRIOR APPLICATION NUMBER: PCT/US03/0408B
PRIOR PILING DATE: 2004-05-24
PRIOR PLING DATE: 2004-04-16
PRIOR PILING DATE: 2004-04-16
PRIOR PILING DATE: 2004-04-16
PRIOR PILING DATE: 2004-01-14
PRIOR PILING DATE: 2003-01-12
PRIOR PILING DATE: 2003-11-23
PRIOR PILING DATE: 2003-12-20
PRIOR PILING DATE: 2003-05-23
PRIOR PILING DATE: 2003-02-20
PRIOR PILING DATE: 2003-05-23
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Fublication No. US20050153916A1

Fublication No. US2005016A1

Fublication No. US2005016A1

Fublication No. US2005016A1

Fublication No. US2005016A1

Fublication No. USA05016A1

Fublication No. USA05017A1

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                                                                                                                                                                                  Sequence 19, Application US/10923330 Publication No. US20050153916A1 GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
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Gaps

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Telomerase Gene (siNA)

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OTHER INFORMATION: Description of Artificial Sequence: Target Sequence/sinh sense re
                                                    APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, James
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RNA INTERFERICE: 400/209 (WABBO2-708-C)
FILE OF INVENTION NUMBER: 9204-08-20
CURRENT FILING DATE: 2004-05-24
PRIOR APPLICATION NUMBER: PCT/US03/0408B
PRIOR FILING DATE: 2004-05-24
PRIOR APPLICATION NUMBER: US 10/826,966
PRIOR PELLING DATE: 2004-01-14
PRIOR FILING DATE: 2004-01-14
PRIOR PELLING DATE: 2004-01-14
PRIOR PELLING DATE: 2003-11-24
PRIOR APPLICATION NUMBER: US 10/757,803
PRIOR PELLING DATE: 2003-11-24
PRIOR PELLING DATE: 2003-11-24
PRIOR PELLING DATE: 2003-11-24
PRIOR PELLING DATE: 2003-12-26
PRIOR PELLING DATE: 2003-12-26
PRIOR PELLING DATE: 2003-12-26
PRIOR PELLING DATE: 2003-12-26
PRIOR PELLING DATE: 2003-05-23
PRIOR PELLING DATE: 2003-05-20
PRIOR PELLING DATE: 2003-05-30
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| APPLICANT: Academic degelumn, Leonid |
| APPLICANT: McSwiggen, James |
| APPLICANT: McSwiggen, James |
| APPLICANT: McSwiggen, Leonid |
| TITLE OF INVENTION: EXPRESSION USING Short Interfering RNA (siNA) |
| TITLE OF INVENTION: EXPRESSION USING Short Interfering RNA (siNA) |
| TITLE OF INVENTION: EXPRESSION USING Short Interfering RNA (siNA) |
| TITLE OF INVENTION: EXPRESSION USING Short Interfering RNA (siNA) |
| TITLE OF INVENTION NUMBER: US 10/923,330 |
| CURRENT FILING DATE: 2004-08-20 |
| PRIOR FILING DATE: 2004-05-24 |
| PRIOR APPLICATION NUMBER: US 60/396,600 |
| PRIOR APPLICATION NUMBER: US 10/826,966 |
| PRIOR FILING DATE: 2004-04-16 |
| PRIOR FILING DATE: 2004-04-16 |
| PRIOR FILING DATE: 2004-04-16 |
| PRIOR PLING DATE: 2004-01-16 |
| PRIOR PLING DATE: 2004-01-14 |
| PRIOR FILING DATE: 2004-01-14 |
| PRIOR FILING DATE: 2003-11-24 |
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                            APPLICANT: Sirna Therapeutics, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; Sequence 24, Application US/10923330; Publication No. US20050153916A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             302 AGAGTTGGGCTCTGTCAGC 320
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 AGAGUUGGGCUCUGUCAGC 19
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SEQ ID NO 23
LENGTH: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: RNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 73.7
Matches 14; Conservative
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                                                                    FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Target Sequence/sinA sense
US-10-923-330-21
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Sequence 22, Application US/1092330

PUBLICALINEORANTION:

APPLICANT: Sirna Therapeutics, Inc.

APPLICANT: McSw4gen, James

FILE REFERENCE: 400/209 (MHH02-708-C)

CURRENT FILING DATE: 2004-08-20

FRICR FILING DATE: 2004-08-20

FRICR FILING DATE: 2004-01-10

FRICR FILING DATE: 2004-01-11

FRICR FILING DATE: 2004-01-14

FRICR FILING DATE: 2004-02-20

FRICR FILING DATE: 2004-02-20

FRICR FILING DATE: 2004-02-20

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FRICR FILING DATE: 2003-03-33

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4.2%; Score 19; DB 1; Length 19;
Best Local Similarity 84.2%; Pred. No. 82;
Matches 16; Conservative 3; Mismatches 0; Indels
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US-10-923-330-23
Sequence 23, Application US/10923330
Publication No. US20050153916A1
GENERAL INFORMATION:
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TYPE: RNA
ORGANISM: Artificial Sequence
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Matches 18; Conserv
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OTHER INFORMATION: Description of Artificial Sequence: Target Sequence/sinA sense re
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APPLICANT: Birna Therapeutics, Inc.
APPLICANT: Birna Therapeutics, Inc.
APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, James
APPLICANT: Beglellan, Leonid
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
TITLE OF INVENTION: RNA INTERFERCE: 2004-08-20
CURRENT FILING DATE: 2004-08-20
FRIOR PAPLICATION NUMBER: US 60/396,600
FRIOR PAPLICATION NUMBER: US 60/396,600
FRIOR APPLICATION NUMBER: US 10/826,966
FRIOR APPLICATION NUMBER: US 10/757,803
FRIOR FILING DATE: 2004-01-16
FRIOR PELING DATE: 2004-01-124
FRIOR PELING DATE: 2003-11-24
FRIOR APPLICATION NUMBER: US 10/757,903
FRIOR PELING DATE: 2003-11-24
FRIOR FILING DATE: 2003-11-24
FRIOR FILING DATE: 2003-11-24
FRIOR FILING DATE: 2003-11-24
FRIOR FILING DATE: 2003-05-20
FRIOR APPLICATION NUMBER: PCT/US03/0508
FRIOR FILING DATE: 2003-05-20
FRIOR FILING DATE: 2003-05-20
FRIOR APPLICATION NUMBER: PCT/US03/0508
FRIOR FILING DATE: 2003-05-20
FRIOR FILING DATE: 2003-05-20
FRIOR APPLICATION NUMBER: PCT/US03/0508
FRIOR FILING DATE: 2003-05-20
FRIOR APPLICATION NUMBER: PCT/US03/0508
FRIOR FILING DATE: 2003-05-20
FRIOR FILING DATE
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Publication No. US20050153916A1
GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: Expression Using Short Interfering RNA (8iNA)
FILE REFERENCE: 400/209 (MBHB02-708-C)
CURRENT APPLICATION NUMBER: US/10/923,330
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82;
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       89.5%; Pred. No. 82;
ive 2; Mismatches
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Best Local Similarity 84.2%; Pred. No. |
Matches 16; Conservative 3; Mismatch
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                                                                                                                          338 CGAGGGGAGGTTCAGGCC 356
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SEQ ID NO 26
LENGTH: 19
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                                               17; Conservative
           Best Local Similarity
                                                                                                                                                                                                                                                                                                                                               JS-10-923-330-26
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US-10-923-330-27
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US-10-923-330-24
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US-10-923-330-25
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US-10-22-310-25
US-10-22-31
US-10-22-32
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PRIOR APPLICATION NUMBER: US 10/693,059
PRIOR PILING DATE: 2003-11-23
PRIOR PLIING DATE: 2003-6-22
PRIOR APPLICATION NUMBER: US 10/444,853
PRIOR APPLICATION NUMBER: PCT/US03/05346
PRIOR PILING DATE: 2003-02-20
PRIOR PLIING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR PILING DATE: 2003-02-20
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEC 1D NOS: 768
SOFTWARE: PatentIn version 3.3
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Pred. No. 82;
3; Mismatches 0; Indels
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: RNA
ORGANISM: Artificial Sequence
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Best Local Similarity
Matches 16; Conserva
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LENGTH: 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FEATURE:
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Score 19; DB 1; Length 19;

4.2%;

Query Match

US09436060A-16.rnpb.sl

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; OTHER INFORMATION: Description of Artificial Sequence: Target Sequence/siNA sense re
US-10-923-330-28
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
APPLICANT: Beigelman, Leoner
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE REFERENCE: 400,209 (MBHB02-708-C)
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT FILING DATE: 2004-08-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
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FRIOR APPLICATION NUMBER: PCT/US03/04088

FRIOR FILING DATE: 2004-05-24

FRIOR FILING DATE: 2002-07-17

FRIOR FILING DATE: 2002-07-17

FRIOR FILING DATE: 2004-05-24

FRIOR FILING DATE: 2004-05-24

FRIOR FILING DATE: 2004-05-24

FRIOR FILING DATE: 2004-05-24

FRIOR PLING DATE: 2004-05-24

FRIOR PLING DATE: 2004-05-34

FRIOR PLING DATE: 2004-05-34

FRIOR PLING DATE: 2004-01-14

FRIOR FILING DATE: 2004-01-14

FRIOR FILING DATE: 2003-11-23

FRIOR FILING DATE: 2003-11-23

FRIOR PLING DATE: 2003-05-23

FRIOR FILING DATE: 2003-05-23

FRIOR FILING DATE: 2003-05-23

FRIOR FILING DATE: 2003-02-20

FRIOR FILING DATE: 2003-02-20

FRIOR PLING DATE: 2003-02-20

FRIOR PLING DATE: 2003-02-20

FRIOR APPLICATION NUMBER: PCT/US03/05028

FRIOR FILING DATE: 2003-02-20

FRIOR FILING DATE: 2003-02-20

FRIOR FILING DATE: 2003-02-20

FRIOR PLING DATE: 2003-02-20

FRIOR PLING DATE: 2003-02-20

FRIOR PLING DATE: 2003-02-20

FRIOR PLING DATE: 2003-02-20

FRIOR APPLICATION NUMBER: PCT/US03/05028

FRIOR PLING DATE: 2003-02-20

FRIOR PLIN
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                                                                     Remaining Prior Application data removed - See File Wrapper or PALM. NUMBER OF SEQ ID NOS: 768
SOFTWARE: Patentin version 3.3
SEQ ID NO 28
LENGTH: 19
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Pred. No. 82;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        392 CGCGCGCGCGCGATTCCC 410
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                                                                                                                                                                                                                                      TYPE: RNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 89.5%;
Matches 17; Conservative
                                         PRIOR FILING DATE: 2003-02-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: RNA
ORGANISM: Artificial Sequence
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Best Local Similarity 78.9
Matches 15; Conservative
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US-10-923-330-28

| Sequence 28, Application US/10923330
| Publication No. US20050153916A1
| GENERAL INFORMATION:
| APPLICANT: Sirra Therapeutics, Inc. |
| APPLICANT: Sirra Therapeutics, Inc. |
| APPLICANT: Bajgelman, Leonid |
| TITLE OF INVENTION: RAN Interference Mediated Inhibition of Telomerase Gene |
| TITLE OF INVENTION: RAN Interference Mediated Inhibition of Telomerase Gene |
| TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA) |
| FILE ERFERENCE: 400/209 (MBHB02-708-C) |
| CURRENT FILING DATE: 2004-08-20 |
| PRIOR PPLICATION NUMBER: US (0/396,600 |
| PRIOR APPLICATION NUMBER: US (0/396,600 |
| PRIOR PLING DATE: 2004-05-24 |
| PRIOR PLING DATE: 2004-05-23 |
| PRIOR PLING DATE: 2003-01-12 |
| PRIOR PLING DATE: 2003-05-23 |
| PRIOR APPLICATION NUMBER: PCT/US03/050-8 |
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Gaps
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4.2%; Score 19; DB 1; Length 19;
Best Local Similarity 94.7%; Pred. No. 82;
Matches 18; Conservative 1; Mismatches 0; Indels
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; OTHER INFORMATION: Description of Artificial Sequence: Target Sequence/sinA sense r*
US-10-923-330-31
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Publication No. US20050153916A1

GENERAL INFORMATION

Publication No. US20050153916A1

GENERAL INFORMATION

GENERAL INFORMATION

FAPLICANT: Michaelgen, James

APPLICANT: Michaelgen, James

APPLICANT: Beigelman, Leonid

TITLE OF INVENTION: Expression Using Short Interfering RNA (sink)

FILE REFERENCE: 400/209 (MBH802-708-C)

CURRENT APPLICATION NUMBER: US/10/923,330

CURRENT FILING DATE: 2004-08-20

FRIOR FILING DATE: 2004-08-20

FRIOR FILING DATE: 2004-07-17

FRIOR FILING DATE: 2004-05-24

FRIOR FILING DATE: 2004-01-14

FRIOR FILIN
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FRIOR FILING DATE: 2004-04-16

FRIOR APPLICATION NUMBER: US 10/757,803

FRIOR FILING DATE: 2004-01-14

FRIOR FILING DATE: 2004-01-14

FRIOR FILING DATE: 2003-01-24

FRIOR FILING DATE: 2003-11-23

FRIOR FILING DATE: 2003-11-23

FRIOR FILING DATE: 2003-05-23

FRIOR PILING DATE: 2003-02-20

FRIOR PAPLICATION NUMBER: PCT/US03/05028

FRIOR PAPLICATION NUMBER: PCT/US03/05028

FRIOR PAPLICATION NUMBER: PCT/US03/05028

FRIOR PILING DATE: 2003-02-20

FRIOR APPLICATION NUMBER: PCT/US03/05028

FRIOR PILING DATE: 2003-02-20

FRIOR PILING DATE: 2003-02-20

FRIOR PILING DATE: 2003-02-20

FRIOR PILING DATE: 2003-02-30

FROM APPLICATION NUMBER: PCT/US03/05028

FROM NUMBER OF SEQ ID NOS: 768

SEQ ID NOS: 768

SEQ ID NOS: 33

FRANCER: 1-1
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4.2%; Score 19; DB 1; Length 19;
Best Local Similarity 84.2%; Pred. No. 82;
Matches 16; Conservative 3; Mismatches 0; Indels
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ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                      APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: Expression Unit Sirna Therefering RNA (sinA)
FILE REPRENCE: 400/209 (MRH020-708-C)
FILE REPRENCE: 400/209 (MRH020-708-C)
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT PELING DATE: 2004-08-20
FRIOR PILING DATE: 2002-07-17
FRIOR FILING DATE: 2002-07-17
FRIOR FILING DATE: 2002-07-17
FRIOR FILING DATE: 2004-05-24
FRIOR FILING DATE: 2004-04-16
FRIOR FILING DATE: 2004-04-16
FRIOR FILING DATE: 2003-11-3
FRIOR FILING DATE: 2003-05-3
FRIOR FILING DATE: 2003-05-3
FRIOR FILING DATE: 2003-05-3
FRIOR FILING DATE: 2003-05-3
FRIOR APPLICATION NUMBER: PCT/US03/05028
FRIOR FILING DATE: 2003-05-3
FRIOR APPLICATION NUMBER: PCT/US03/05028
FRIOR APPLICATION TO PALICATION 3.3
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4.2%; Score 19; DB 1; Length 19;
Best Local Similarity 89.5%; Pred. No. 82;
Matches 17; Conservative 2; Mismatches 0; Indels
                                                                                                                                                                Sequence 30, Application US/10923330 Publication No. US20050153916A1 GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 103
US-10-923-330-31
                                                                                                                              US-10-923-330-30
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PUBLICANT: Sitna Therapeutics, Inc.
GENERAL INFORMATION:
TO SEQUENCE 39, Application US/1092330

PUBLICANT: Sitna Therapeutics, Inc.
APPLICANT: McSwigen, James
APPLICANT: Bigelman, Leonid
TITLE OF INVENTION: Expression Using Short Interfering RNA (sink)
FILE REPRENCE: 400/209 (MHHB02-708-C)
FILE REPRENCE: 400/209 (MHHB02-708-C)
FILE REPRINCE: 400/209 (MHHB02-708-C)
FILE REPRINCE: 2004-08-20
FRIOR PRILING DATE: 2004-08-20
FRIOR PRILING DATE: 2004-08-20
FRIOR PRILING DATE: 2004-08-24
FRIOR FILING DATE: 2004-01-14
FRIOR FILING DATE: 2004-01-16
FRIOR FILING DATE: 2003-11-23
FRIOR FILING DATE: 2003-11-23
FRIOR APPLICATION NUMBER: US 10/693,059
FRIOR FILING DATE: 2003-02-20
FRIOR FILING DATE: 2003-02-20
FRIOR FILING DATE: 2003-02-20
FRIOR FILING DATE: 2003-03-65-23
FRIOR FILING DATE: 2003-02-20
FRIOR FILING DATE: 2003-03-30
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                                   FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region US-10-923-330-38
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                                                                                                                                                                                      Query Match
4.2%; Score 19; DB 1; Length 19;
Best Local Similarity 100.0%; Pred. No. 82;
Matches 19; Conservative 0; Mismatches 0; Indels
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4.2%; Score 19; DB 1; Length 19;
Best Local Similarity 100.0%; Pred. No. 82;
Matches 19; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                       14 TGGGCCTGGGAGGGGTGGT 32
                                                                                                                                                                                                                                                                                                                                                                                         19 TGGGCCTGGGAGGGGTGGT 1
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ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 105
US-10-923-330-39/c
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US-10-923-330-40/c
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RESULT 108

US-10-323-330-42/c

| Sequence 42, Application US/1022330
| Publication No. US20050153916A1
| GENERAL INFORMATION:
| APPLICANT: Sirrar Therappentics, Inc.
| APPLICANT: Beigelman, Leonid
| TITLE OF INVENTION: NRM Interference Mediated Inhibition of Telomerase Gene
| TITLE OF INVENTION: NRM Interference Mediated Inhibition of Telomerase Gene
| TITLE OF INVENTION: NRM Interference Mediated Inhibition of Telomerase Gene
| TITLE OF INVENTION: NRM Interference Mediated Inhibition of Telomerase Gene
| TITLE OF INVENTION: NUMBER: US0/203 (MRH802-008-C)
| CURRENT APPLICATION NUMBER: US 60/396,600
| PRIOR FILING DATE: 2004-05-24
| PRIOR PELICATION NUMBER: US 60/396,600
| PRIOR FILING DATE: 2004-05-24
| PRIOR APPLICATION NUMBER: US 10/720,448
| PRIOR APPLICATION NUMBER: US 10/720,448
| PRIOR PELING DATE: 2004-01-124
| PRIOR PELING DATE: 2003-11-24
| PRIOR PELING DATE: 2003-11-24
| PRIOR PELING DATE: 2003-11-24
| PRIOR PELING DATE: 2003-11-23
| PRIOR PELING DATE: PELING DATE: PELING DATE: PELING DATE: PELING DATE: PELING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region US-10-923-330-41
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PRIOR FILING DATE: 2003-11-23
PRIOR APPLICATION NUMBER: US 10/444,853
PRIOR FILING DATE: 2003-05-23
PRIOR FILING DATE: 2003-05-20
PRIOR FILING DATE: 2003-02-20
PRIOR PLICATION NUMBER: PCT/US03/05028
PRIOR FILING DATE: 2003-02-20
PRIOR FILING DATE: 2003-02-20
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 768
SOFTHARE: PATENTIN 9-10 NOS: 768
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100.0%; Pred. No. 82;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      68 TAGGCGCCGTGCTTTTGCT 86
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: RNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 100.(
Matches 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                   SEQ ID NO 41
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4.2%; Score 19; DB 1; Length 19; 100.0%; Pred. No. 82;

Query Match Best Local Similarity

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GENERAL INFORMATION:
APPLICANT: BITA THEAPPENTION:
APPLICANT: BITA THEAPPENTION:
APPLICANT: BITA THEAPPENTION:
APPLICANT: BEGGEDAMA: Lecond
TITLE OF INVENTION: RAM Interference Mediated Inhibition of Telomerage Gene
TITLE OF INVENTION: RAM Interference Mediated Inhibition of Telomerage Gene
TITLE OF INVENTION: RAM Interference Mediated Interfering RNA (siNA)
TITLE OF INVENTION: RAM INTERFERENCE: 400/209 (MRHBD2.708-C)
CURRENT FILING DATE: 2004-08-20
FRIOR PAPLICATION NUMBER: US/10/930,600
FRIOR PELING DATE: 2004-05-24
FRIOR APPLICATION NUMBER: US 80/396,600
FRIOR APPLICATION NUMBER: US 10/757,803
FRIOR PILING DATE: 2004-01-14
FRIOR PILING DATE: 2004-01-14
FRIOR PELING DATE: 2004-01-14
FRIOR PELICATION NUMBER: US 10/757,803
FRIOR APPLICATION NUMBER: US 10/757,803
FRIOR PELING DATE: 2003-11-23
FRIOR PELING DATE: 2003-11-24
FRIOR PELING DATE: 2003-11-24
FRIOR PELING DATE: 2003-05-23
FRIOR PELING DATE: 2003-05-23
FRIOR PELING DATE: 2003-05-20
FRIOR APPLICATION NUMBER: US 10/444,853
FRIOR PELING DATE: 2003-05-20
FRIOR PELING DATE: 2003-05-20
FRIOR PELING DATE: 2003-05-20
FRIOR FILING DATE: 2003-05-30
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Publication No. US20050153916A1
GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Beigelman, Leonid
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT FILING DATE: 2004-08-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gape
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                                                                                                                                                                                                                                                                                     Sequence 43, Application US/10923330 Publication No. US20050153916A1 GENERAL INFORMATION:
86 TCCCCGCGCGCTGTTTTC 104
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                                            19 TCCCCGCGCGCTGTTTTC 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                             -10-923-330-43/c
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; OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region US-10-923-330-45
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US-10-923-330-46
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Remaining Prior Application data removed - See File Wrapper or PALM. NUMBER OF SEQ ID NOS: 768 SOFTWARE: PatentIn version 3.3 SEQ ID NO 45
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; Sequence 46, Application US/10923330
; Publication No. US20050153916A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                         140 GCCGCCTTCCACCGTTCAT 158
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              158 TTCTAGAGCAAACAAAAA 176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19 GCCGCCTTCCACCGTTCAT 1
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                                                                                                                                                             TYPE: RNA
ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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i US-10-923-3310-45/c
i Sequence 45, Application US/10923330
publication No. US20050153916A1
i GENERAL INFORMATION:
    APPLICANT: Sirna Therapeutics, Inc.
    APPLICANT: Beigelman, Leonie
    TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
    TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
    TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
    TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
    TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
    TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
    CURRENT APPLICATION NUMBER: US/10/923,330
    CURRENT PILING DATE: 2004-08-20
    PRIOR FILING DATE: 2004-05-24
    PRIOR FILING DATE: 2004-05-24
    PRIOR FILING DATE: 2004-05-24
    PRIOR PILING DATE: 2004-05-24
    PRIOR FILING DATE: 2004-01-16
    PRIOR FILING DATE: 2003-11-24
    PRIOR FILING DATE: 2003-11-24
    PRIOR FILING DATE: 2003-11-23
    PRIOR FILING DATE: 2003-11-23
    PRIOR FILING DATE: 2003-11-23
    PRIOR FILING DATE: 2003-11-23
    PRIOR PILING DATE: 2003-01-12-3
    PRIOR RELING DATE: 2003-01-13-3
    PRIOR RELING DATE: 2003-02-20
    PRIOR PILING DATE: 2003-02-20
    PRIOR PILING DATE: 2003-02-20
    PRIOR PILING DATE: 2003-02-20
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                                        ## FRIOR APPLICATION NUMBER: US 60/396,600
#PRIOR FILING DATE: 2002-07-17
#PRIOR FILING DATE: 2002-07-17
#PRIOR FILING DATE: 2004-05-24
#PRIOR PELING DATE: 2004-05-05
#PRIOR PELING DATE: 2004-05-05
#PRIOR PELING DATE: 2004-04-16
#PRIOR APPLICATION NUMBER: US 10/757,803
#PRIOR PILING DATE: 2004-01-14
#PRIOR PILING DATE: 2003-11-24
#PRIOR PILING DATE: 2003-11-24
#PRIOR PILING DATE: 2003-11-23
#PRIOR APPLICATION NUMBER: US 10/444,853
#PRIOR APPLICATION NUMBER: PC1/US03/05946
#PRIOR APPLICATION NUMBER: PC1/US03/05346
#PRIOR PILING DATE: 2003-02-23
#PRIOR PILING DATE: 2003-02-20
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   APPLICATION NUMBER: PCT/US03/04088
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        122 GCGGAAAGCCTCGGCCTG 140
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ORGANISM: Artificial Sequence
                                     2004-05-24
                              FILING DATE: 2004-09
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local S:
Matches 19
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JS-10-923-330-49/c
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        US.10-233-330-470.
US.10-233-330-470.
US.0020153916A1
Sequence 47, Application US/10923330
Publication No. US20050153916A1
CERNEAL INFORMATION.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Games
APPLICANT: Games
APPLICANT: Games
APPLICANT: Games
APPLICANT: Games
APPLICANT: MARSAGGAN, James
APPLICANT: MARSAGGAN, James
TITLE OF INVENTION: RAPECATION NUMBER: US/10/923,330
CURRENT FLING DATE: 2004-05-24
PRIOR PELING DATE: 2004-05-24
PRIOR APPLICATION NUMBER: PCT/US04/16390
PRIOR PELING DATE: 2004-05-24
PRIOR APPLICATION NUMBER: US 60/396,600
PRIOR APPLICATION NUMBER: US 60/396,600
PRIOR APPLICATION NUMBER: US 10/757,803
PRIOR PELING DATE: 2004-05-24
PRIOR APPLICATION NUMBER: US 10/757,803
PRIOR PELING DATE: 2004-01-14
PRIOR PELING DATE: 2003-01-17
PRIOR PELING DATE: 2003-01-12
PRIOR PELING DATE: 2003-01-14
PRIOR PELING DATE: 2003-01-14
PRIOR PELING DATE: 2003-01-12
PRIOR PELING DATE: 2003-01-13
PRIOR PELING DATE: 2003-
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APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, James
APPLICANT: Beigelman, Leonies
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE REPERENCE: 400/229 (MBH802-708-C)
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT FILING DATE: 2004-08-20
PRIOR APPLICATION NUMBER: PCT/US03/04088
PRIOR FILING DATE: 2004-05-24
PRIOR FILING DATE: 2004-05-24
PRIOR FILING DATE: 2004-05-24
PRIOR FILING DATE: 2002-07-17
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PRIOR FILING DATE: 2004-05-24
PRIOR APPLICATION NUMBER: US 10/826,966
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-330-47/c
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APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Beigelman, Jemes
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE REFERENCE: 400/209 (MBHB02-708-C)
CURRENT APPLICATION NUMBER: BCT/US03/04088
PRIOR FILING DATE: 2004-05-24
PRIOR PELING DATE: 2004-05-24
PRIOR FILING DATE: 2004-05-24
PRIOR FILING DATE: 2004-06-24-16
PRIOR FILING DATE: 2004-06-24-16
PRIOR FILING DATE: 2004-04-16
PRIOR FILING DATE: 2004-01-14
PRIOR FILING DATE: 2003-11-24
PRIOR FILING DATE: 2003-11-23
PRIOR PLILING DATE: 2003-05-33
PRIOR PLILING DATE: 2003-05-33
PRIOR PLILING DATE: 2003-05-33
PRIOR PLILING DATE: 2003-05-34
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US-10-923-330-48
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82;
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LING DATE: 2004-04-16
PLICATION NUMBER: US 10/757,803
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ORGANISM: Artificial Sequence
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| Publication No. US20050153916A1
| GENERAL INFORMATION:
| APPLICANT: Sitna Therapeutics, Inc.
| APPLICANT: Sitna Therapeutics, Inc.
| APPLICANT: McSwiggen, James
| APPLICANT: McSwiggen, James
| APPLICANT: Beigelman, Leonid
| TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
| TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
| FILE REPRENCE: 400/209 (MRHB02-708-C)
| CURRENT PEPLICATION NUMBER: US/10/923,330
| FURENT PELING DATE: 2004-08-20
| PRIOR PAPLICATION NUMBER: US 0/396,600
| PRIOR PELING DATE: 2004-01-14
| PRIOR FILING DATE: 2004-11-24
| PRIOR FILING DATE: 2004-11-24
| PRIOR FILING DATE: 2003-11-24
| PRIOR FILING DATE: 2003-11-23
| PRIOR FILING DATE: 2003-11-33
| PRIOR FILING DATE: 2003-10-30
| PRIOR FILING DATE
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region
US-10-923-330-49
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4.2%; Score 19; DB 1; Length 19;
Best Local Similarity 100.0%; Pred. No. 82;
Matches 19; Conservative 0; Mismatches 0; Indels
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US-10-923-330-51/c
; Sequence 51, Application US/10923330
; Publication No. US2050153916A1
; GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
; APPLICANT: McSwiggen, James
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ORGANISM: Artificial Sequence
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| APPLICANT: Basquaran, Looned | APPLICANT: Basquaran, Basqu
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JS-10-923-330-55/c
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Publication No. US20050153916A1

Sequence 53, Application US/10923330

Publication No. US20050153916A1

APPLICANT: Missing Therapeutics, Inc.

APPLICANT: Missing Therapeutics, Inc.

APPLICANT: Missing No. US20050153916A1

TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene

TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene

TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene

TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene

TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene

TITLE OF INVENTION: RNA INTERF. US/10/203, 330

CURRENT PELLOR DATE: 2004-06-24

PRIOR PELLOR DATE: 2004-06-24

PRIOR PELLOR DATE: 2004-06-24

PRIOR APPLICATION NUMBER: US 10/826,966

PRIOR FILING DATE: 2004-01-14

PRIOR PILING DATE: 2004-01-14

PRIOR PILING DATE: 2003-01-14

PRIOR PILING DATE: 2003-01-14

PRIOR PILING DATE: 2003-01-14

PRIOR PELLOR DATE: 2003-01-14

PRIOR PELLOR DATE: 2003-01-14

PRIOR APPLICATION NUMBER: US 10/693,059

PRIOR PELLOR DATE: 2003-01-12

PRIOR PELLOR DATE: 2003-02-20

PRIOR PELLOR PELLOR DATE: 2003-02-20

PRIOR PELLOR DATE: 200
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US-10-923-330-53
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Pred. No. 82;
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PRIOR APPLICATION NUMBER: US 10/444,853
PRIOR FILING DATE: 2003-05-23
PRIOR PILING DATE: 2003-02-20
PRIOR PILING DATE: 2003-02-20
PRIOR PILING DATE: 2003-02-20
PRIOR PILING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR PILING DATE: 2003-02-20
NUMBER OF SEQ ID NOS: 768
SOFTWARE: PATENTIN VERSION 3.3
LENGTH: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     266 CCGGGCTTCTCCGGAGGC 284
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ORGANISM: Artificial Sequence
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4.2%; Score 19; DB 1; Length 19; 100.0%; Pred. No. 82; tive 0; Mismatches 0; Indels

Query Match
Best Local Similarity 100.
Matches 19; Conservative

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; OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region US-10-923-330-54
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Sequence 55, Application US/10923330
Publication No. US20050153916A1
GENERAL INFORMATION:
APPLICANT: Sirna Therapeutice, Inc.
APPLICANT: Sirna Therapeutice, Inc.
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: Rapression Using Short Interfering RNA (81NA)
FILE REFERENCE: 400/209 (MBHB02-708-C)
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT FILING DATE: 2004-08-20
PRIOR APPLICATION NUMBER: PCT/US03/04088
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                                                                                              Sequence 54, Application US/10923330 Publication No. US20050153916A1 GENERAL INFORMATION:
284 CACCCACTGCCACCGCGAA 302
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        302 AGAGTTGGGCTCTGTCAGC 320
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Gaps

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; OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region US-10-923-330-56
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                                                                                                                                                                                                                                                                                                                                                               Length 19;
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82;
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Best Local Similarity 100.0%; Pred. No. 82;
Matches 19; Conservative 0; Mismatches
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; Publication No. US20050153916A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19 CGAGGCGAGGTTCAGGCC 1
                                                       SOFTWARE: PatentIn version 3.3
SEQ ID NO 56
LENGTH: 19
                                                                                                                                                     TYPE: RNA
ORGANISM: Artificial Sequence
FEATURE:
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ORGANISM: Artificial Sequence
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Best Local Similarity
Matches 19; Conserva
                           NUMBER OF SEQ ID NOS:
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US-10-923-330-57/c
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APPLICANT: SITMA Thexagelulics, Inc.
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RNA PERSONAL THEORY (SINA)
FILE REFERENCE: 400/209 (MBHB02-708-C)
CURRENT PELLING DATE: 2004-08-20
PRIOR PRICATION NUMBER: PCT/US03/04088
PRIOR APPLICATION NUMBER: DCT/US03/04088
PRIOR FILING DATE: 2004-05-24
PRIOR PELLING DATE: 2004-05-24
PRIOR PELLING DATE: 2004-06-24
PRIOR PELLING DATE: 2004-06-24
PRIOR PELLING DATE: 2004-06-24
PRIOR PELLING DATE: 2004-06-24
PRIOR PELLING DATE: 2004-06-14
PRIOR PELLING DATE: 2004-01-14
PRIOR PELLING DATE: 2004-01-14
PRIOR PELLING DATE: 2003-11-23
PRIOR PELLING DATE: 2003-12-23
PRIOR PELLING DATE: 2003-05-23
PRIOR PELLING DATE: 2003-02-20
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PRIOR FILING DATE: 2004-05-24

PRIOR PLICATION NUMBER: US 60/396,600

PRIOR FILING DATE: 2002-07-17

PRIOR FILING DATE: 2004-05-24

PRIOR FILING DATE: 2004-05-24

PRIOR PLICATION NUMBER: US 10/826,966

PRIOR PELING DATE: 2004-04-16

PRIOR PLING DATE: 2004-01-14

PRIOR PLING DATE: 2004-01-14

PRIOR PLING DATE: 2003-11-24

PRIOR PLING DATE: 2003-11-24

PRIOR PLING DATE: 2003-11-23

PRIOR PLING DATE: 2003-11-23

PRIOR PLING DATE: 2003-11-23

PRIOR PLING DATE: 2003-10-20

PRIOR PLING DATE: 2003-10-20

PRIOR PLING DATE: 2003-05-20

PRIOR PLING DATE: 2003-05-20

PRIOR PLING DATE: 2003-02-20

PRIOR PLING D
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APPLICANT: Sirna Therapeutics, Inc.
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ORGANISM: Artificial Sequence
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APPLICATION NUMBER: US 10/757,803
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ORGANISM: Artificial Sequence
FEATURE:
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US-10-923-330-59/c

i Sequence 59, Application US/10923330

i Publication No. US20050153916A1

i GENERAL INFORMATION:

APPLICANT: Sirna Therapeutica, Inc.

APPLICANT: Beigelman, Leonid

TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene

TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)

FILE REFERENCE: 400/209 (MBHB02-708-C)

CURRENT FILING DATE: 2004-08-20

PRIOR PLICATION NUMBER: PCT/US03/04088

PRIOR PLICATION NUMBER: BC 60/396,600

PRIOR PLICATION NUMBER: US 60/396,600

PRIOR PLICATION NUMBER: US 60/396,600

PRIOR PLICATION NUMBER: US 60/396,600

PRIOR PLILING DATE: 2004-05-24

PRIOR PLILING DATE: 2004-05-24

PRIOR FILING DATE: 2004-05-24

PRIOR FILING DATE: 2004-05-24

PRIOR FILING DATE: 2004-05-34

PRIOR FILING DATE: 2004-05-34
                                                                                                                                                                                APPLICANT: SITRIA inTerapeutrs, int.

APPLICANT: Baigelman, Leonid

APPLICANT: Beigelman, Leonid

TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)

TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)

FILE REPRENCE: 400/209 (MBHB02-708-C)

CURRENT APPLICATION NUMBER: US/10/923,330

CURRENT APPLICATION NUMBER: US/10/923,330

PRIOR APPLICATION NUMBER: US 60/396,600

PRIOR PLING DATE: 2004-06-20

PRIOR PLING DATE: 2004-06-21

PRIOR PLING DATE: 2004-06-24

PRIOR PLING DATE: 2004-06-24

PRIOR PLING DATE: 2004-01-14

PRIOR PLING DATE: 2004-01-14

PRIOR PLING DATE: 2004-01-14

PRIOR PLING DATE: 2004-01-14

PRIOR PLING DATE: 2003-01-1-24

PRIOR PLING DATE: 2003-01-24

PRIOR PLING DATE: 2003-01-23

PRIOR PLING DATE: 2003-01-23

PRIOR PLING DATE: 2003-02-20

PRIOR PLING DATE: 2003-02-3

PRIOR PLING DATE: 2003-02-4

PRIOR PLING DATE: 2003-02-4

PRIOR PLING DATE: 2003-02-4

PRIOR PLING DATE: 2003-02-4

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                                      Sequence 58, Application US/10923330; Publication No. US20050153916A1; GENERAL INFORMATION: APPLICANT: Sirna Herapeutics, Inc.
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ORGANISM: Artificial Seguence
        US-10-923-330-58/c
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PRDICARIN NO. USES COUGOLISTICANT.

APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSWiggen, James
APPLICANT: Beigelman, Lection
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Corresponding Manager, 100/209 (MBHB02-708-C)
FILE REFERENCE: 400/209 (MBHB02-708-C)
CURRENT FILING DATE: 2004-00-20
FRIOR FILING DATE: 2004-01-17
FRIOR FILING DATE: 2004-01-52
FRIOR FILING DATE: 2004-01-64-16
FRIOR FILING DATE: 2004-01-63
FRIOR FILING DATE: 2004-01-63
FRIOR FILING DATE: 2004-01-23
FRIOR FILING DATE: 2003-01-23
FRIOR FILING DATE: 2003-01-23
FRIOR FILING DATE: 2003-01-23
FRIOR FILING DATE: 2003-01-20
FRIOR FILING DATE: 2003-02-20
FRIOR FILING DATE: 2003-03-20-20
FRIOR FILING DATE: 2
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                                           PRIOR PELICATION NUMBER: US 10/720,448
PRIOR FILING DATE: 2003-11-24
PRIOR PLLING DATE: 2003-11-24
PRIOR PLLING DATE: 2003-11-24
PRIOR PLLING DATE: 2003-11-23
PRIOR PLLING DATE: 2003-01-23
PRIOR PLLING DATE: 2003-05-23
PRIOR PLICATION NUMBER: PCT/US03/05346
PRIOR PLICATION NUMBER: PCT/US03/05346
PRIOR FILING DATE: 2003-02-20
PRIOR FIL
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FILING DATE: 2004-01-14
APPLICATION NUMBER: US 10/720,448
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; OTHER INFORMATION: oligonucleotide US-10-831-266-6
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ORGANISM: Artificial Sequence
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Matches 19; Conservative
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UG-10-921-330-61/C

Sequence 61, Application US/10923330

Publication No. US2005013916A1

SEGNERAL INFORMATION:

APPLICANT: MCGRANTION:

APPLICANT: MCGRANTION:

APPLICANT: MCGRANTION:

TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene

TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)

TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)

TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)

TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)

FILE REPRENCE: 400/209 (HRHBOZ-708-C)

CURRENT PELLORICATION NUMBER: US/04-08-C)

FRIOR APPLICATION NUMBER: US/04-08-C)

FRIOR APPLICATION NUMBER: US 10/826,600

FRIOR APPLICATION NUMBER: US 10/826,966

FRIOR APPLICATION NUMBER: US 10/757,803

FRIOR FILING DATE: 2004-01-14

FRIOR FILING DATE: 2003-11-24

FRIOR FILING DATE: 2003-11-24

FRIOR FILING DATE: 2003-11-24

FRIOR PELLORITON NUMBER: US 10/693,059

FRIOR APPLICATION NUMBER: US 10/20-346

FRIOR FILING DATE: 2003-02-20

FRIOR APPLICATION NUMBER: US 10/20-346

FRIOR FILING DATE: 2003-02-20

FRIOR APPLICATION NUMBER: US 10/20-366

FRIOR FILING DATE: 2003-02-20

FRIOR FILING DATE: 2003-03-20-20

FRIOR FILING DATE: 2003-03-20-20
OTHER INFORMATION: Description of Artificial Sequence: siNA antisense region
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                                                                                                        DB 1; Length 19;
82;
                                                                                                                                                                              0; Indels
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4.2%; Score 19; DB 1; Length 19;
Best Local Similarity 100.0%; Pred. No. 82;
Matches 19; Conservative 0; Mismatches 0; Indels
                                                                                              Query Match 4.2%; Score 19; DB Best Local Similarity 100.0%; Pred. No. 82, Matches 19; Conservative 0; Mismatches
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Publication No. US20050153916A1
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APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
APPLICANT: Beigelman, Leonid
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          ; OTHER INFORM
US-10-923-330-60
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LENGTH: 19
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TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE REPERENCE: 400/209 (MHHBD2-708-C)
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT APPLICATION NUMBER: US/10/923,330
FRIOR FILING DATE: 2004-08-24
FRIOR FILING DATE: 2004-05-24
FRIOR FILING DATE: 2004-05-24
FRIOR FILING DATE: 2004-05-24
FRIOR FILING DATE: 2004-05-24
FRIOR FILING DATE: 2004-05-34
FRIOR FILING DATE: 2004-01-124
FRIOR PELING DATE: 2004-01-124
FRIOR PELING DATE: 2003-11-24
FRIOR PELING DATE: 2003-11-24
FRIOR PELING DATE: 2003-11-23
FRIOR PELING DATE: 2003-05-33
FRIOR FILING DATE: 2003-05-33
FRIOR FIL
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US-10-923-330-62
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100.0%; Pred. No. 82;
Live 0; Mismatches 0; Indels
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| Sequence 6, Application US/10831266
| Publication No. US200500340431
| GENERAL INFORMATION:
| APPLICANT: ROWLey, Peter T.
| TITLE OF INVENTION: TELOMERASE INTERFERENCE
| FILE REFERENCE: A-71506-1/RFT/PHR
| CURRENT PEDLICATION NUMBER: US/10/831,266
| PRIOR PAPLICATION NUMBER: US/10/831,266
| PRIOR FILING DATE: 2002-10-16
| PRIOR FILING DATE: 2001-10-22
| PRIOR FILING DATE: 2001-10-22
| PRIOR PELING DATE: 2002-02-10
| PRIOR APPLICATION NUMBER: US 60/359,196
| PRIOR FILING DATE: 2002-02-20
| PRIOR FILING DATE: 2002-02-20
| PRIOR FILING DATE: 2002-05-22
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Gaps
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                          Query Match 4.2%; Score 19; DB 1; Length 21; Best Local Similarity 100.0%; Pred. No. 1e+02; Matches 19; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                               Sequence 9, Application US/10831266
| Publication No. US2005003404A1
| GENERAL INFORMATION:
| APPLICANT: ROWLEY, Peter T.
| TITLE OF INVENTION: TELOMERASE INTERFERENCE TILE REFERENCE A-71506-1/RFT/THR
| CURRENT APPLICATION NUMBER: US/10/831,266
| CURRENT FILING DATE: 2004-04-22/RPIOR FILING DATE: 2001-10-16
| PRIOR APPLICATION NUMBER: US 60/345,326
| PRIOR APPLICATION NUMBER: US 60/359,196
| PRIOR FILING DATE: 2002-02-20
| PRIOR PELING DATE: 2002-02-20
| PRIOR PELING DATE: 2002-05-22
| NUMBER OF SEQ ID NOS: 17
| SOFTWARE: PALENTIN VERSION 3.2
| SEQ ID NO 9
| LENGTH: 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 6, Application US/10831267
| Publication No. US2005009177A1
| GENERAL INFORMATION:
| APPLICANT: ROWLEY, Peter T.
| TITLE OF INVENTION: TELOMERASE INTERFERENCE FILE REFRENCES A-71506-2/RFT/THR
| CURRENT APPLICATION NUMBER: US/10/831,267 CURRENT FILING DATE: 2004-04-22 PRIOR APPLICATION NUMBER: US/0/33146 PRIOR FILING DATE: 2001-10-16 PRIOR APPLICATION NUMBER: US 60/345,326 PRIOR APPLICATION NUMBER: US 60/345,326 PRIOR APPLICATION NUMBER: US 60/359,196 PRIOR APPLICATION NUMBER: US 60/359,196 PRIOR PRIOR DATE: 2002-02-20 PRIOR APPLICATION NUMBER: US 60/383,195 PRIOR PRILING DATE: 2002-05-22 NUMBER OF SEQ ID NOS: 23 SOFTWARE: PALENT UN VERSION 3.2 SEQ ID NO S: SEQ ID NOS: 23 LENGTH: 21
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FEATURE:
OTHER INFORMATION: oligonucleotide
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                                                                                                                                             270 GGCTTCTCCGGAGGCACCC 288
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  Score 19; DB 1; Length 21; Pred. No. 1e+02;
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4.2%; Score 19; DB 1; Length 21;
Best Local Similarity 100.0%; Pred. No. 1e+02;
Matches 19; Conservative 0; Mismatches 0; Indels
                                                           0; Indels
                                                                                                                                                                                                                                                                         Sequence 7, Application US/10831266

Sequence 7, Application US/10831266

Publication No. US200S0003404A1

GENERAL INFORMATION:
TITLE OF INVENTION: TELEOMERASE INTERFERENCE
FILE REFERENCE: A-71506-1/RRT/THR

CURRENT APPLICATION NUMBER: US/10/831,266

CURRENT APPLICATION NUMBER: US/10/831,266

PRIOR APPLICATION NUMBER: US/00333065

PRIOR FILING DATE: 2002-10-16

PRIOR FILING DATE: 2001-10-22

PRIOR APPLICATION NUMBER: US 60/345,326

PRIOR FILING DATE: 2001-10-22

PRIOR APPLICATION NUMBER: US 60/345,326

PRIOR FILING DATE: 2001-10-22

PRIOR PILING DATE: 2002-02-20

PRIOR FILING DATE: 2002-02-20

PRIOR FILING DATE: 2002-02-22

NUMBER OF SEQ ID NOS: 17

SEQ ID NO 7

LENGTH: 21
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Sequence 8, Application US/10831266

Sequence 8, Application US/10831266

GENERAL INPORMATION:
TITLE OF INVENTION: TELCOMERASE INTERFERENCE
FILE REFERENCE: A-71506-1/RET/THR
CURRENT APPLICATION NUMBER: US/10/831,266

CURRENT FILING DATE: 2004-04-22

PRIOR APPLICATION NUMBER: US/10/831,266

PRIOR APPLICATION NUMBER: US 60/345,326

PRIOR FILING DATE: 2002-10-16

PRIOR FILING DATE: 2001-10-22

PRIOR FILING DATE: 2001-10

PRIOR FILING DATE: 2001-02

PRIOR PLICATION NUMBER: US 60/385,196

PRIOR FILING DATE: 2002-02-20

PRIOR PLING DATE: 2002-02-20

PRIOR FILING DATE: 2002-02-20

PRIOR FILING DATE: 2002-02-20

SOFTWARE OF SEQ ID NOS: 17

SOFTWARE: PARCHALLY NUMBER: US 60/383,195

PRIOR FILING DATE: 2002-05-22

NUMBER OF SEQ ID NOS: 17

SEQ ID NO 8

LENGTH: 21
Query Match 4.2%; Score 19; DB Best Local Similarity 68.4%; Pred. No. 1e+0 Matches 13; Conservative 6; Mismatches
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; OTHER INFORMATION: oligonucleotide
US-10-831-266-8
                                                                                                                   42 TTGTCTAACCCTAACTGAG 60
                                                                                                                                                1 UUGUCUAACCCUAACUGAG 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 42 TIGICIAACCCIAACIGAG 60
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ORGANISM: Artificial
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ORGANISM: Artificial
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US-10-831-267-9/c
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US-10-923-330-546
                                                                                          Query Match
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                                                    Score 19; DB 1; Length 21; Pred. No. 1e+02;
                                                                                                    0; Indels
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100.0%; Pred. No. 16+02;
Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                  Sequence 7, Application US/10811267
Sedence 7, Application US/10811267
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: ROWLEY, Peter T.
TITLE OF INVENTION: TELOMERASE INTERFERENCE
FILE REFERENCE: A-7156-2/RFT/THR
CURRENT FILING DATE: 2004-04-22
FRIOR APPLICATION NUMBER: US/10/831,267
CURRENT FILING DATE: 2002-10-16
PRIOR APPLICATION NUMBER: US 60/345,326
PRIOR FILING DATE: 2001-10-22
PRIOR PELING DATE: 2001-10-22
PRIOR APPLICATION NUMBER: US 60/359,196
PRIOR FILING DATE: 2002-02-0
PRIOR APPLICATION NUMBER: US 60/389,195
PRIOR PILING DATE: 2002-05-2
PRIOR APPLICATION NUMBER: US 60/383,195
PRIOR FILING DATE: 2002-05-2
SEQ ID NOS: 23
SOFTWARE: PALENTIN VERSION 3.2
SEQ ID NO 7
LENGTH: 21
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Sequence 8, Application US/10831267

Sequence 8, Application US/10831267

GENERAL INFORMATION:

TITLE OF INVENTION: TELCMERARE INTERFERENCE

FILE REFERENCE: A-71506-2/RFT/THR

CURRENT APPLICATION NUMBER: US/10/831,267

CURRENT APPLICATION NUMBER: US/10/831,267

CURRENT APPLICATION NUMBER: US/03/33146

PRIOR APPLICATION NUMBER: US 60/345,326

PRIOR FILING DATE: 2001-10-16

PRIOR FILING DATE: 2001-10-2

PRIOR APPLICATION NUMBER: US 60/345,326

PRIOR FILING DATE: 2001-10-2

PRIOR APPLICATION NUMBER: US 60/359,196

PRIOR APPLICATION NUMBER: US 60/383,195

PRIOR RILING DATE: 2002-02-2

PRIOR APPLICATION NUMBER: US 60/383,195

PRIOR ETLING DATE: 2002-02-2

NUMBER OF SEQ ID NOS: 23

SOFTWARE: PALENTING DATE: 2002-05-22

NUMBER OF SEQ ID NOS: 23

SEQ ID NO 8

LENGTH:: 21
                                                                                                 6; Mismatches
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                                                                                                                                            42 TTGTCTAACCCTAACTGAG 60
                                                                                                                                                                       1 UNGUCUAACCCUAACUGAG 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               42 TIGICIAACCCIAACTGAG 60
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                                          Query Match 4.2%;
Best Local Similarity 68.4%;
Matches 13; Conservative (
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Best Local Similarity
Matches 19; Conserva
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ORGANISM: Artificial
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US-10-831-267-6
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Sequence 546, Application US/10923330

Publication No. US20050153916A1

GENERAL INFORMATION:
APPLICANT: Birna Therapeutics, Inc.
APPLICANT: McSwiggen, James
APPLICANT: Basigelman, Leonid
ITILE OF INVENTION: RAY Interference Mediated Inhibition of Telomerase Gene
ITILE OF INVENTION: RAY INTERFERENCE: A00/209
CURRENT APPLICATION NUMBER: US/10/203,330
CURRENT APPLICATION NUMBER: PCT/US03/04088
PRIOR PILING DATE: 2004-08-20
PRIOR PLING DATE: 2004-05-24
PRIOR PLING DATE: 2004-05-16
PRIOR FLING DATE: 2004-05-16
PRIOR FLING DATE: 2004-05-16
PRIOR FLING DATE: 2004-04-16
PRIOR FLING DATE: 2004-04-16
PRIOR FLING DATE: 2004-01-14
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                                                                                            DB 1; Length 21; 1e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                Sequence 9, Application US/10831267
Sequence 9, Application US/10831267
GENERAL INFORMATION:
APPLICANT: ROWLEY, Peter T.
TITLE OF INVENTION: TELOMERASE INTERFERENCE
FILE REFERENCE: A-7156-2/RFT/THR
CURRENT APPLICATION NUMBER: US/10/831,267
CURRENT FILING DATE: 2004-04-22
FRIOR APPLICATION NUMBER: US/10/831,267
FRIOR APPLICATION NUMBER: US 60/345,326
FRIOR FILING DATE: 2001-10-22
FRIOR FILING DATE: 2001-10-22
FRIOR APPLICATION NUMBER: US 60/359,196
FRIOR FILING DATE: 2002-02-2
FRIOR APPLICATION NUMBER: US 60/383,195
FRIOR FILING DATE: 2002-05-22
NUMBER OF SEQ ID NOS: 23
SEQ ID NO 9
LENGTH: 21
LENGTH: 21
                                                                                      Query Match
4.2%; Score 19; DB
Best Local Similarity 100.0%; Pred. No. 10+
Matches 19; Conservative 0; Mismatches
; OTHER INFORMATION: oligonucleotide US-10-831-267-8
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                                                                                                                                                                                                         270 GGCTTCTCCGGAGGCACCC 288
                                                                                                                                                                                                                                              1 GGCTTCTCCGGAGGCACCC 19
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ORGANISM: Artificial
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FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: sinA sense region
                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 63.2%;
Matches 12; Conservative
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| Sequence 547, Application US/20050153916A1
| Sequence 547, Application US20050153916A1
| CENERAL INFORMATION:
| APPLICANT: Sina Therapeutics, Inc.
| APPLICANT: McSwiggen, James
| APPLICANT: Beigelman, Leonid
| TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
| FILE REFRENCE: 400/209 (MRHEN2-708-C)
| FILE REFRENCE: 400/209 (MRHEN2-708-C)
| FILE REPRENCE: 400/209 (MRHEN2-708-C)
| FILING DATE: 2004-08-20
| PRIOR APPLICATION NUMBER: US 60/396,600
| PRIOR PILING DATE: 2004-06-24
| PRIOR FILING DATE: 2004-06-14
| PRIOR FILING DATE: 2004-06-14
| PRIOR FILING DATE: 2004-06-16
| PRIOR FILING DATE: 2004-06-16
| PRIOR FILING DATE: 2004-06-16
| PRIOR FILING DATE: 2004-01-14
| PRIOR PILING DATE: 2003-01-14
| PRIOR PILING DATE: 2003-11-3
| PRIOR PILING DATE: 2003-11-3
| PRIOR PILING DATE: 2003-11-3
| PRIOR FILING DATE: 2003-11-3
| PRIOR FILING DATE: 2003-11-3
| PRIOR FILING DATE: 2003-11-3
| PRIOR PILING DATE: 2003-05-20
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PRIOR PILING DATE: 2003-11-24
PRIOR APPLICATION NUMBER: US 10/693,059
PRIOR PILING DATE: 2003-11-23
PRIOR PELING DATE: 2003-01-23
PRIOR APPLICATION NUMBER: PCT/US03/05346
PRIOR PILING DATE: 2003-02-20
PRIOR PILING DATE: 2003-02-20
PRIOR PILING DATE: 2003-02-20
PRIOR PILING DATE: 2003-02-20
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 768
SOFTMARE: PATENTIN Version 3.3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: misc_feature

// LOCATION: (20)...(21)

OTHER INFORMATION: n stands for thymidine

US-10-923-330-546
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NUMBER OF SEQ ID NOS: 768
SOFTWARE: PatentIn version 3.3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: RNA
ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                SEQ ID NO 546
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APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Bacgelman, Leonid
TITLE OF INVENTION: Bacgelman, Leonid
TITLE OF INVENTION: Expression Using Short Interfering RNA (sINA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (sINA)
FILE REPERENCE: 400/209 (MBHBO2-708-C)
FILE REPERENCE: 400/209 (MBHBO2-708-C)
FILE REPERENCE: 2004-08-20
FRIOR PELICATION NUMBER: US/10/923,330
CURRENT APPLICATION NUMBER: US 60/396,600
FRIOR FILING DATE: 2004-05-24
FRIOR FILING DATE: 2004-06-24
FRIOR FILING DATE: 2004-06-24
FRIOR FILING DATE: 2004-06-24
FRIOR FILING DATE: 2004-01-14
FRIOR FILING DATE: 2004-01-14
FRIOR FILING DATE: 2004-11-24
FRIOR FILING DATE: 2003-11-24
FRIOR FILING DATE: 2003-12-23
FRIOR FILING DATE: 2003-12-24
FRIOR FILING DATE: 2003-12-24
FRIOR FILING DATE: 2003-02-20
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SOFTWARE: Patentin version 3.3
SEQ ID NO 548
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Pred. No. 1e+02;
7; Mismatches
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4.2%; Score 19; DB 1;
Best Local Similarity 68.4%; Pred. No. 1e+02;
Matches 13; Conservative 6; Mismatches
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NAMME/KEY: misc feature
LOCATION: (20) -. (21)
OTHER INFORMATION: n stands for thymidine
US-10-923-330-548
FEATURE:
NAME/KEY: misc_feature
COCATION: (20)...(21)
OTHER INFORMATION: n stands for thymidine
US-10-923-330-547
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                    Sequence 549, Application US/1092330
| Fouricria 2015-549 |
| Sequence 549, Application US/1092330 |
| Publication No. US20050153916A1 |
| APPLICANT: Sirna Therapeutics, Inc. |
| APPLICANT: McKWiggen, James |
| TITLE OF INVENTION: Expression Using Short Interfering RNA (SiNA) |
| FILE REPERENCE | 2004-08-20 |
| FRICR PELICATION NUMBER: US/10/923,330 |
| CURRENT APPLICATION NUMBER: US 0/306,600 |
| PRIOR PELICATION NUMBER: US 0/04-04-16 |
| PRIOR PELICATION NUMBER: US 10/720,448 |
| PRIOR PLICATION NUMBER: US 10/303-02-03 |
| PRIOR PLICATION NUMBER: US 10/303-03-05-03 |
| PRIOR PLICATION NUMBER: US 10/303-03-05-03 |
| PRIOR PLICATION NUMBER: US 10/303-02-03 |
| PRIOR PLICATION NUMBER: US 10/303-02-03 |
| PRIOR PLICATION NUMBER: US 10/303-02-03 |
| PRIOR PLICATION NUMBER: US 2003-02-20 |
| PRIOR PLICATION NUMBER: US 2003-02-20 |
| PRIOR PLICATION NUMBER: US 2003-02-20 |
| PRIOR PLICATION NUMBER: US 2003-02-03 |
| PRIOR PLICATION DATE: 2003-02-03 |
| PRIOR PLICATION NUMBER: US 2003-02-03 |
| PRIOR PLICATION DE APPLICATION NUMBER: US 2003-02-03 |
| PRIOR PLICATION NUMBER: US 2003-02-03 |
| PRIOR PLICATION DE APPLICATION DE APP
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APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: RAY Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE REFERENCE: 400/209 (MBHB02-708-C)
CURRENT APPLICATION NUMBER: US/10/923, 330
CURRENT FILING DATE: 2004-08-20
PRIOR PILING DATE: 2004-08-24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: misc feature
LOCATION: (20)...(21)
CTHER INFORMATION: n stands for thymidine
US-10-923-330-549
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ORGANISM: Artificial Sequence
US-10-923-330-549
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US-10-923-330-550
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PRIOR PRICACION NAMES: 18 6/356,600

PRIOR PLICATION NAMES: 1002-07-17

PRIOR PLICATION NAMES: 1002-07-17

PRIOR PLICATION NAMES: 1004-01-17

PRIOR PLICATION NAMES: 1004-01-12

PRIOR PLICATION NAMES: 1004-01-12

PRIOR PLICATION NAMES: 1004-01-12

PRIOR PLICATION NAMES: 1004-01-12

PRIOR PLICATION NAMES: 1007-01-12

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15 Sequence 552, Application US/10923330

15 Publication No. US20050153916A1

1 Publication No. US20050153916A1

APPLICANT: WESViggen, James

APPLICANT: MESViggen, James

APPLICANT: MESViggen, Jeonid

TITLE OF INVENTION: RAN Interference Mediated Inhibition of Telomerase Gene

TITLE OF INVENTION: RAN AINEARE WESTORS (MBHB02-708-C)

CURRENT APPLICATION NUMBER: US/10/923,330

CURRENT FILING DATE: 2004-08-24

PRIOR FILING DATE: 2004-08-24

PRIOR PILING DATE: 2004-05-24

PRIOR PILING DATE: 2004-05-24

PRIOR PILING DATE: 2004-01-14

PRIOR PILING DATE: 2004-01-14

PRIOR PILING DATE: 2004-01-14

PRIOR PILING DATE: 2004-01-14

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                                                                                  - See File Wrapper or PALM.
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Pred. No. 1e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                            ; NAME/KEY: misc_feature
; LOCATION: (20)..(21)
; OTHER INFORMATION: n stands for thymidine
US-10-923-330-551
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// LCCATION: (20)...(21)

// OTHER INFORMATION: n stands for thymidine

US-10-923-330-552
PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR FILING DATE: 2003-02-20
Remaining Prior Application data removed
NUMBER OF SEQ ID NOS: 768
SOFTWARE: Patentin version 3.3
SEQ ID NO 551
LENGTH: 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 73.7%;
Matches 14; Conservative
                                                                                                                                                                                                                                                                      TYPE: RNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Artificial Sequence
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US-10-923-330-552
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OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region
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USE-10-923-3130-554/C
USE-10-923-3130-554/C
USE-10-923-3130-554/C
USE-10-923-3130-554/C
USE-10-923-3130-554/C
USEDICANT: Sitna Therapeutics, Inc.
APPLICANT: McGswiggen, James
APPLICANT: McGswiggen, James
APPLICANT: McSwiggen, Jeneid
TITLE OF INVENTION: Expression using Short Interfering RNA (siNA)
TITLE OF INVENTION: Expression using Short Interfering RNA (siNA)
TITLE OF INVENTION: Expression using Short Interfering RNA (siNA)
TITLE OF INVENTION: WHERE: US/10/923,330
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT APPLICATION NUMBER: US (0/396,600)
FRIOR FILING DATE: 2004-05-24
FRIOR PELING DATE: 2004-05-24
FRIOR APPLICATION NUMBER: US 10/405-966
FRIOR PELING DATE: 2004-04-16
FRIOR APPLICATION NUMBER: US 10/750,448
FRIOR APPLICATION NUMBER: US 10/750,448
FRIOR APPLICATION NUMBER: US 10/693,059
FRIOR APPLICATION NUMBER: US 10/693,059
FRIOR APPLICATION NUMBER: US 10/693,059
FRIOR PELING DATE: 2003-11-23
FRIOR APPLICATION NUMBER: US 10/444,853
FRIOR APPLICATION NUMBER: PCT/US03/05346
FRIOR PELING DATE: 2003-05-23
FRIOR PELING DATE: 2003-05-23
FRIOR PELING DATE: 2003-05-24
FRIOR PELING DATE: 2003-05-23
FRIOR PELING DATE: 2003-05-24
FRIOR PELING DATE: 2003-05-26
FRIOR PELING DATE: 2003-05-36
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Length 21;
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4.2%; Score 19; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 1e+02;
Matches 19; Conservative 0; Mismatches 0;
; DB 1;
. le+02;
                                                                     5; Mismatches
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Score 19;
Pred. No. 1
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GENERAL INFORMATION: APPLICANT: Sirna Therapeutics, Inc.
                                                                                                                                           300 GAAGAGTTGGGCTCTGTCA 318
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  145 CTTCCACCGTTCATTCTAG 163
                                                                                                                                                                                       1 GAAGAGUUGGGCUCUGUCA 19
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SEQ ID NO 554
LENGTH: 21
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Query Match
Best Local Similarity 73.7%;
Matches 14; Conservative
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US-10-923-330-555/c
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147 TCCACCGTTCATTCTAGAG 165
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ORGANISM: Artificial Sequence
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NAME/KEY: misc feature
LOCATION: (20)...(21)
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APPLICANT: McSwiggen, James
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: RAY Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RAYDERSON USING Short Interfering RNA (siNA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE REFERENCE: 400/209 (MBHB02-708-C)
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT PILING DATE: 2004-08-20
PRIOR PILING DATE: 2004-08-20
PRIOR PLICATION NUMBER: PCT/US04/16390
PRIOR PLICATION NUMBER: US 60/396,600
PRIOR PLICATION NUMBER: US 10/826,966
PRIOR PLICATION NUMBER: US 10/826,966
PRIOR PLICATION NUMBER: US 10/757,803
PRIOR PLING DATE: 2004-01-14
PRIOR PLING DATE: 2004-01-16
PRIOR PLING DATE: 2003-01-23
PRIOR PLING DATE: 2003-11-23
PRIOR PLING DATE: 2003-11-23
PRIOR PLING DATE: 2003-11-23
PRIOR PLING DATE: 2003-02-20
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Publication No. US20050153916A1

GENERAL INFORMATION:
APPLICANT: SITUA Therapeutics, Inc.
APPLICANT: Beigellan, James
APPLICANT: Beigellan, Leonid
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
CURRENT APPLICATION NUMBER: US/10/923,330
PRIOR APPLICATION NUMBER: US 60/396,600
PRIOR APPLICATION NUMBER: US 60/396,600
PRIOR APPLICATION NUMBER: US 10/4-05-24
PRIOR FILING DATE: 2004-05-24
PRIOR FILING DATE: 2004-05-24
PRIOR FILING DATE: 2004-05-24
PRIOR FILING DATE: 2004-05-16
PRIOR FILING DATE: 2004-04-16
PRIOR FILING DATE: 2004-04-16
PRIOR FILING DATE: US 10/757,803
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SOFTWARE: PatentIn version 3.3
SEQ ID NO 555
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COTHER INFORMATION: n stands for thymidine US-10-923-330-555
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ORGANISM: Artificial Sequence
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FRICH PILLING DATE; 2004-01-14

FRICH APPLICATION NUMBE; US 10/750,448

FRICH APPLICATION NUMBE; US 10/750,448

FRICH APPLICATION NUMBE; US 10/453,059

FRICH APPLICATION NUMBE; US 10/44.633

FRICH APPLICATION NUMBE; US 10/44.633

FRICH APPLICATION NUMBE; DETUGNOS/346

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LOCATION: (20)..(21)
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APPLICANT: SITMS Theragelulicity, Inc.
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RNA INTERFERENCE: 400/209 (MBHB02-708-C)
CURRENT PAPPLICATION NUMBER: US/10/923,330
CURRENT PAPLICATION NUMBER: DCT/US03/0408
PRIOR APPLICATION NUMBER: DCT/US03/0408
PRIOR PILING DATE: 2002-07-17
PRIOR PILING DATE: 2004-05-24
PRIOR PILING DATE: 2004-06-24
PRIOR PILING DATE: 2004-04-16
PRIOR PILING DATE: 2004-04-16
PRIOR PILING DATE: 2004-04-16
PRIOR PILING DATE: 2003-11-23
PRIOR PILING DATE: 2003-11-23
PRIOR PILING DATE: 2003-11-23
PRIOR PILING DATE: 2003-05-23
                                                                                             OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region
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NUMBER OF SEQ ID NOS: 768
SOFTWARE: PatentIn version 3.3
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Pred. No. 1e+02;
0; Mismatches 0; Indels
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4.2%; Score 19; DB 1; Length 21;
Best Local Similarity 100.0%; Pred. No. 1e+02;
Matches 19; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                      Query Match 4.2%; Score 19; DB Best Local Similarity 100.0%; Pred. No. 164 Matches 19; Conservative 0; Mismatches
                                                                                                                                                                                          LOCATION: (20) (21)
COTHER INFORMATION: n stands for thymidine US-10-923-330-557
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) LOCATION: (20)...(21)
) OTHER INFORMATION: n stands for thymidine
US-10-923-330-558
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; Sequence 558, Application US/10923330; Publication No. US20050153916A1; GENERAL INFORMATION: APPLICANT: Sirna Therapeutics, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     148 CCACCGTTCATTCTAGAGC 166
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TYPE: RNA
ORGANISM: Artificial Sequence
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                                                                                                                           FEATURE:
NAME/KEY: misc_feature
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LENGTH: 21
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Page 58

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FEATURE:
NAME/KEY: misc_feature
LOCATION: (20)...(20)
OTHER INFORMATION: n stands for thymidine
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NAME/KEY: misc_feature
LOCATION: (7)...(8)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
FEATURE:
NAME/KEY: misc_feature
LOCATION: (10)...(12)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
FEATURE:
FEATURE:
FEATURE:
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NAME/KEY: misc_feature
LOCATION: (1)..(5)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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Best Local Similarity
Matches 12; Conserva
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                                                                                                                                                                                                                                                                                                                                TYPE: RNA
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Sequence 562, Application US/10923330

Publication No. US20050153916A1

GENERAL INFORMATION:

APPLICANT: Sirna Therapeutics, Inc.

APPLICANT: Beigelman, Leonid

TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene

TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase of TITLE OF INVENTION: RAY INTERFERENCE: 400/209 (MBHB02-708-C)

TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)

FILE REFERENCE: 400/209 (MBHB02-708-C)

CURRENT APPLICATION NUMBER: US/10/923,330

CURRENT APPLICATION NUMBER: PCT/US03/04088

PRIOR FILING DATE: 2004-08-20

PRIOR FILING DATE: 2004-05-24

PRIOR FILING DATE: 2004-05-24

PRIOR FILING DATE: 2004-01-14

PRIOR FILING DATE: 2004-01-14

PRIOR APPLICATION NUMBER: US 10/757,803

PRIOR FILING DATE: 2004-01-14

PRIOR FILING DATE: 2003-11-24

PRIOR FILING DATE: 2003-11-24

PRIOR FILING DATE: 2003-11-23

PRIOR PRIOR FILING DATE: 2003-11-23
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                                                                           FRIOR FILING DATE: 2004-05-17

FRIOR FILING DATE: 2004-05-24

FRIOR FILING DATE: 2004-05-24

FRIOR FILING DATE: 2004-05-16

FRIOR FILING DATE: 2004-05-16

FRIOR FILING DATE: 2004-04-16

FRIOR FILING DATE: 2004-04-14

FRIOR FILING DATE: 2004-01-14

FRIOR FILING DATE: 2003-11-24

FRIOR FILING DATE: 2003-11-24

FRIOR FILING DATE: 2003-11-24

FRIOR FILING DATE: 2003-11-24

FRIOR FILING DATE: 2003-12-24

FRIOR PLING DATE: 2003-12-24

FRIOR PLING DATE: 2003-12-24

FRIOR PAPLICATION NUMBER: US 10/693,059

FRIOR APPLICATION NUMBER: US 10/444,853

FRIOR APPLICATION NUMBER: PT/US03/05346

FRIOR PILING DATE: 2003-02-20

FRIOR FILING DATE: 2003-02-30

FRIOR FILING DATE: 2003-02-30
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4.2%; Score 19; DB 1; Length 21;
Best Local Similarity 100.0%; Pred. No. 1e+02;
Matches 19; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; LOCATION: (20)-.(21); OTHER INFORMATION: n stands for thymidine US-10-923-330-560
APPLICATION NUMBER: PCT/US03/04088
                                FILING DATE: 2004-05-24
APPLICATION NUMBER: US 60/396,600
FILING DATE: 2002-07-17
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Sequence 563, Application US/10923330

Publication No. US20050153916A1

GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE REPERBACE: 400/209 (MHBD2-708-C)
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT APPLICATION NUMBER: US/10/923,330
FRIOR APPLICATION NUMBER: DCT/US03/04088
FRIOR APPLICATION NUMBER: DCT/US03/04088
FRIOR PILING DATE: 2004-05-24
FRIOR PILING DATE: 2002-07-17
FRIOR PILING DATE: 2002-07-17
FRIOR APPLICATION NUMBER: PCT/US04/16390
FRIOR APPLICATION NUMBER: DCT/US04/16390
FRIOR APPLICATION NUMBER: US/10/9204
FRIOR APPLICATION NUMBER: US/10/9204
FRIOR APPLICATION NUMBER: US/10/9204
FRIOR PILING DATE: 2004-05-24
FRIOR PILING DATE: 2004-06-4-16
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PRIOR APPLICATION NUMBER: PCT/US03/05346
PRIOR FILING DATE: 2003-02-20
PRIOR FILING DATE: 2003-02-20
PRIOR FILING DATE: 2000-02-20
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 768
SOFTWARE: Patentin version 3.3
SEGO ID NO 562
LENGTH: 21
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LOCATION: (1). 7(1)
OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
NAME/KEY: misc feature
LOCATION: (21) - (21)
COTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity
US-10-923-330-562
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4.2%; Score 19; DB 1; Length 21; 63.2%; Pred. No. 1e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      7; Mismatches
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PCT/US03/04088

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RESULT 154
US-10-923-330-565
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US-10-923-330-564
Sequence 564, Application US/10923330
Publication No. US20050153916A1
SERNERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE REPERENCE: 400/209 (MBHB02-708-C)
CURRENT APPLICATION NUMBER: US/10/923, 330
CURRENT FILING DATE: 2004-08-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: Description of Artificial Sequence: siNA sense region
PRIOR APPLICATION NUMBER: US 10/757,803
PRIOR PLILING DATE: 2004-01-14
PRIOR PLILING DATE: 2003-11-24
PRIOR PLILING DATE: 2003-11-23
PRIOR PLILING DATE: 2003-11-23
PRIOR PLILING DATE: 2003-11-23
PRIOR PLILING DATE: 2003-05-23
PRIOR PLILING DATE: 2003-05-23
PRIOR PLILING DATE: 2003-05-20
PRIOR PLILING DATE: 2003-02-20
PRIOR PRIOR PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR PLILING DATE: 2003-02-20
PRIOR PLING DATE: 2003-02-20
PRIOR PLING DATE: 2003-02-20
PRIOR PLING DATE: 2003-02-30
PRIOR APPLICATION NUMBER: PCT/US03/05048
PRIOR PLING DATE: 2003-02-30
PRIOR P
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LOCATION: (1)._(1)
OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               // NAME/KEY: misc_feature
// LOGATION: (21)...(21)
// OTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity
US-10-923-330-563
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 4.2%; Score 19; DB 1; Length 21; Best Local Similarity 63.2%; Pred. No. 1e+02; Matches 12; Conservative 7; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY: misc_feature
LOCATION: (20) ... (20)
OTHER INFORMATION: n stands for thymidine
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LOCATION: (9)._(11)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (13) ... (16)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (1). 7.(4)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (6)._(7)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 UUCCACCGUUCAUUCUAGA 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: RNA
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FRIOR APPLICATION UNDERS: US 60/396,600
FRIOR FILING DATE: 2002-07-17
FRIOR APPLICATION NUMBER: US 60/396,600
FRIOR FILING DATE: 2002-07-17
FRIOR FILING DATE: 2004-05-24
FRIOR FILING DATE: 2004-05-24
FRIOR FILING DATE: 2004-01-46
FRIOR FILING DATE: 2004-01-41
FRIOR FILING DATE: 2004-01-41
FRIOR FILING DATE: 2003-11-24
FRIOR PELICATION NUMBER: US 10/720,448
FRIOR PELICATION NUMBER: US 10/693,059
FRIOR FILING DATE: 2003-11-23
FRIOR FILING DATE: 2003-11-23
FRIOR FILING DATE: 2003-11-23
FRIOR FILING DATE: 2003-05-23
FRIOR FILING DATE: 2003-05-23
FRIOR FILING DATE: 2003-02-20
FRIOR FILING DATE: 2003-02-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LOCATION: (1). 7(1)
OTHER INFORMATION: 5'-3 attached terminal deoxyabasic mosity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: Description of Artificial Sequence:
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OTHER INFORMATION: n stands for thymidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: misc feature
LOCATION: (1). (3)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LOCATION: (8). (10)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LOCATION: (5). (6)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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; Publication No. US20050153916A1
; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: RNA ORGANISM: Artificial Sequence
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LOCATION: (12)..(15)
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Indels

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14; Conservative
                             Matches
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                        APPLICANT: Beigelman, Leonid

TITLE OF INVENTION: Experence Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Experence Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Experence Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Experence Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Experence Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Experence 100/209 (WBHB02-708-C)
CURRENT APPLICATION WUMBER: US/10/923,330
CURRENT PILING DATE: 2004-06-24
PRIOR PELING DATE: 2004-05-24
PRIOR PELING DATE: 2004-05-24
PRIOR APPLICATION WUMBER: US 10/826,966
PRIOR PELING DATE: 2004-01-14
PRIOR PELING DATE: 2004-01-14
PRIOR PELING DATE: 2004-01-14
PRIOR PELING DATE: 2003-01-124
PRIOR PELING DATE: 2003-11-24
PRIOR PELING DATE: 2003-11-24
PRIOR PELING DATE: 2003-11-24
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PRIOR PELING DATE: 2003-05-23
PRIOR PELING DATE: 2003-05-23
PRIOR PELING DATE: 2003-05-23
PRIOR PELING DATE: 2003-05-23
PRIOR PELING DATE: 2003-02-20
PRIOR PELING DATE: 2003-02-20
PRIOR PELING DATE: 2003-02-3
PRIOR 
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NAME/KEY: misc_feature
LOCATION: (1)...(1)
OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
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. OTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity US-10-923-330-565
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OTHER INFORMATION: n stands for thymidine
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LOCATION: (1)._(2)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (4)._(5)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (11) ... (14)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (19)...(19)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
APPLICANT: Sirna Therapeutics, Inc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: RNA
ORGANISM: Artificial Sequence
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LOCATION: (7)..(9)
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NAME/KEY: misc feature
LOCATION: (20)..(20)
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LOCATION: (21)...(21)
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Length 21;

Score 19; DB 1; Pred. No. 1e+02;

4.2%;

Query Match Best Local Similarity

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Oy 148 COCCCUCANCICACACC 19

RESULT 185

I CCACCCUCANUCACACC 185

I PUBLICANT INTERCEDENT 187

I RESPERENT INTERCEDENT 187

I RESPECT 1
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FEATURE:
NAME/KEY: misc_feature
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APPLICANT: Baigelman, Leonid
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE OF INVENTION: 2004-05-24
FRIOR APPLICATION NUMBER: US 60/396,600
FRIOR FILING DATE: 2004-05-24
FRIOR PELING DATE: 2004-05-24
FRIOR PELING DATE: 2004-05-24
FRIOR PELING DATE: 2004-05-34
FRIOR PELING DATE: 2004-01-14
FRIOR PELING DATE: 2004-01-14
FRIOR PELING DATE: 2003-11-23
FRIOR PELING DATE: 2003-11-23
FRIOR PELING DATE: 2003-11-23
FRIOR PELING DATE: 2003-11-23
FRIOR PELING DATE: 2003-05-23
FRIOR APPLICATION NUMBER: US 10/444,853
FRIOR PELING DATE: 2003-02-20
FRIOR APPLICATION NUMBER: US 10/444,853
FRIOR PELING DATE: 2003-02-20
FRIOR PELING DATE: 2003-02-20
FRIOR PELING DATE: 2003-02-20
FRIOR APPLICATION NUMBER: PET/US03/05028
FRIOR FILING DATE: 2003-02-20
FRIOR PELING DATE: 2003-03-20
FRIOR PELING DATE: 2003-03-20
FRIOR PELING DATE: 200
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                           NAME/KEY: misc_feature

1 LOCATION: (21)...(21)

2 CHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity

US-10-923-330-566
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NAME/KEY: misc_feature
LOCATION: (1)..(1)
OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
                                                                                                                                                                                                                                                                                                                                                                              Query Match
4.2%; Score 19; DB 1; Length 21;
Best Local Similarity 73.7%; Pred. No. 1e+02;
Matches 14; Conservative 5; Mismatches 0; Indels
       FEATURE:
NAME/KEY: misc_feature
LOCATION: (20) .. (20)
OTHER INFORMATION: n stands for thymidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: misc feature
LOCATION: (5). 77
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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NAME/KEX: misc_feature
LOCATION: (2). (3)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             , Sequence 567, Application US/10923330
, Publication No. US20050153916A1
, GENERAL INFORMATION:
, APPLICANT: Sirna Therapeutics, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              149 CACCGTTCATTCTAGAGCA 167
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: RNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 156
US-10-923-330-567
                                                                                                                                                           FEATURE:
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GENERAL INFORMATION The Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggan, James
APPLICANT: McSwiggan, James
APPLICANT: McSwiggan, James
APPLICANT: McSwiggan, James
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE REFERENCE: 400/209 (WHBH02-708-C)
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT PILING DATE: 2004-06-20
FRIOR PRIOR PLING DATE: 2004-06-20
FRIOR PILING DATE: 2004-07-17
FRIOR PILING DATE: 2004-07-17
FRIOR FILING DATE: 2004-04-16
FRIOR FILING DATE: 2004-05-23
FRIOR FILING DATE: 2004-05-23
FRIOR FILING DATE: 2003-05-23
FRIOR FILING DATE: 2003-05-24
FRIOR FILING DATE: 2003-05-34
FRIOR FILING DATE: 200
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
4.2%; Score 19; DB 1; Length 21;
Best Local Similarity 73.7%; Pred. No. 1e+02;
Matches 14; Conservative 5; Mismatches 0; Indels
                                                                           FEATURE:
NAME/KEY: misc_feature
NAME/KEY: misc_feature
CCCATION: (17)...(17)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
FEATURE:
NAME/KEY: misc_feature
LOCATION: (20)...(20)
OTHER INFORMATION: n stands for thymidine
LOCATION: (9). (12)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 568, Application US/10923330 Publication No. US20050153916A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     150 ACCGTTCATTCTAGAGCAA 168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 ACCGUUCAUUCUAGAGCAA 19
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: misc_feature
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LOCATION: (1)..(1)
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## STATUS APPLICANT NO. USZOUGOLES 916A1

| GENERAL INFORMATION:
| APPLICANT: Sirna Therapeutics, Inc.
| APPLICANT: Sirna Therapeutics, Inc.
| APPLICANT: Mosaiggen, James
| TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
| TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
| TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
| TITLE OF INVENTION: RNA INTERF. US/10/923,330
| CURRENT FILING DATE: 2004-08-24
| PRIOR FILING DATE: 2004-06-24
| PRIOR FILING DATE: 2004-01-16
| PRIOR PELICATION NUMBER: US 10/32,466
| PRIOR PELICATION NUMBER: US 10/720,448
| PRIOR PELICATION NUMBER: US 10/720,448
| PRIOR FILING DATE: 2004-01-14
| PRIOR FILING DATE: 2003-11-24
| PRIOR FILING DATE: 2003-11-24
| PRIOR FILING DATE: 2003-11-23
| PRIOR PELICATION NUMBER: US 10/693,059
| PRIOR PELICATION NUMBER: US 10/693,059
| PRIOR FILING DATE: 2003-02-20
| PRIOR PELICATION NUMBER: PCT/US03/05028
| P
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COATION: (20)...(20)

OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
FEATURE:
NAME/KEY: misc_feature

LOCATION: (21)...(21)

COTHER INFORMATION: n stands for thymidine
US-10-923-330-570
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
4.2%; Score 19; DB 1;
Best Local Similarity 100.0%; Pred. No. 1e+02;
Matches 19; Conservative 0; Mismatches (
                                                                                                                                                                          NAME/KEY: misc feature
LOCATION: (7)...(7)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                            DOCATION: (1)..(2)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                 FEATURE:
NAME/KEY: misc feature
LOCATION: (11) -. (11)
FEATURE:
FEATURE:
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LOCATION: (14)...(14)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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Publication No. US20050153916A1
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                         misc_feature
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Sequence 570, Application US/10923330

Fablication No US20050153916A1

GENERALI INFORMATION:
APDLICAMY: Star Therapeutics, Inc.
APDLICAMY: Birar Therapeutics, Inc.
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
CURRENT APDLICATION NUMBER: 904-06-20

PRIOR FILING DATE: 2004-06-24
PRIOR FILING DATE: 2004-05-24
PRIOR PELLOR NUMBER: US 10/826, 966
PRIOR FILING DATE: 2004-05-11
PRIOR PELLOR OF APDLICATION NUMBER: US 10/720, 448
PRIOR PELLOR OF APDLICATION NUMBER: US 10/720, 448
PRIOR PELLOR DATE: 2004-01-11-23
PRIOR PELLOR DATE: 2004-01-11-23
PRIOR PELLOR DATE: 2004-01-11-23
PRIOR PELLOR DATE: 2003-05-23
PRIOR DATE: 2003-05-23
PRIOR DATE: 2003-05-23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
4.2%; Score 19; DB 1; Length 21;
Best Local Similarity 73.7%; Pred. No. 1e+02;
Matches 14; Conservative 5; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LOCATION: (20)...(20)
OTHER INFORMATION: n stands for thymidine
NAME/KEY: misc_feature
LOCATION: (7)..(8)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                     NAME/KEY: misc feature
LOCATION: (12)...(15)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                            FEATURE:
NAME/KRY: misc feature
LOCATION: (17). (18)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: misc_feature
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Gaps

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Length 21; 0; Indels

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Sequence 373, Application US/10923330

**Publication No. US20050153916A1

**Publication No. US20050153916A1

**Generication No. US20050153916A1

**GENERAL INFORMATION:

**APPLICANT: Sirna Therapeutice, Inc.

**APPLICANT: Beigelman, Leonid

**ITILE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene

**ITILE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene

**ITILE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene

**ITILE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene

**ITILE OF INVENTION: RNABER: US/10/923,330

**CURRENT APPLICATION NUMBER: US/10/923,330

**CURRENT FILING DATE: 2004-09-20

**PRIOR PILING DATE: 2004-09-20

**PRIOR PILING DATE: 2004-09-16

**PRIOR PILING DATE: 2004-09-16

**PRIOR PILING DATE: 2004-09-16

**PRIOR APPLICATION NUMBER: US 10/757,803

**PRIOR APPLICATION NUMBER: US 10/757,803

**PRIOR PILING DATE: 2003-11-24

**PRIOR PILING DATE: 2003-11-24

**PRIOR PILING DATE: 2003-11-24

**PRIOR PILING DATE: 2003-11-23

**PRIOR PILING DATE: 2003-11-23
                                                                                                                                                                                                                                                                INFORMATION: Description of Artificial Sequence: sinh antisense region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: misc feature
LOCATION: (20)...(20)
OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 4.2%; Score 19; DB 1; Length 21; Best Local Similarity 100.0%; Pred. No. 1e+02; Matches 19; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COTATION: (21). (21) COTHER INFORMATION: n stands for thymidine US-10-923-330-572
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PRIOR FILING DATE: 2003-05-23
PRIOR APPLICATION NUMBER: PCT/US03/05346
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FEATURE:
NAME/KEY: misc_feature
LOCATION: (16)..(16)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY: misc_feature
LOCATION: (9)._(9)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LOCATION: (13)...(13)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                      NAME/KEY: misc_feature
LOCATION: (1)...(4)
OCHER INFORMATION: 2'-deoxy-2'-fluoro
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          147 TCCACCGTTCATTCTAGAG 165
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19 rccaccerrcarrcragae 1
NUMBER OF SEQ ID NOS: 768
SOFTWARE: Patentin version 3.3
SEQ ID NO 572
LENGTH: 21
                                                                                                                                                                                          ORGANISM: Artificial Sequence
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LOCATION: (21)..(21)
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US-10-923-330-573/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE:
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APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
APPLICANT: Belgelman, Leonid
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE REPERENCE: 2004-08-20
CURRENT APPLICATION NUMBER: DC/10/923,330
CURRENT FILING DATE: 2004-08-20
PRIOR PELING DATE: 2004-05-24
PRIOR FILING DATE: 2004-05-24
PRIOR FILING DATE: 2004-05-24
PRIOR PELING DATE: 2004-01-14
PRIOR FILING DATE: 2004-01-14
PRIOR FILING DATE: 2004-01-14
PRIOR FILING DATE: 2003-01-12
PRIOR FILING DATE: 2003-11-23
PRIOR FILING DATE: 2003-11-23
PRIOR FILING DATE: 2003-11-23
PRIOR FILING DATE: 2003-02-23
PRIOR FILING DATE: 2003-02-23
PRIOR PELING DATE: 2003-02-23
PRIOR PELING DATE: 2003-02-23
PRIOR APPLICATION NUMBER: US 10/444,853
PRIOR PELING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR PELING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR PRIOR PRIOR DATE: 2003-02-20
PRIOR PRIOR PRIOR PRIOR DATE: 2003-02-20
PRIOR PRIOR PRIOR DATE: 2003-02-20
PRIOR PRIOR PRIOR DATE: 2003-02-20
PRIOR PRIOR PRIOR PRIOR DATE: 2003-02-20
                                                                                                                    OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region
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OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 4.2%; Score 19; DB 1; Length 21; Best Local Similarity 100.0%; Pred. No. 1e+02; Matches 19; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LOCATION: (21)...(21)
CTHER INFORMATION: n stands for thymidine US-10-923-330-571
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY: misc feature
LOCATION: (12)...(12)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: misc_feature
LOCATION: (15)...(15)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                         LOCATION: (1)...(3)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                   FEATURE:
NAME/KEY: misc_feature
NAME/KEY: wisc_feature
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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; Sequence 572, Application US/10923330
; Publication No. US20050153916A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          146 TTCCACCGTTCATTCTAGA 164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19 rrccaccerrcarreraga 1
                                          ORGANISM: Artificial Sequence
                                                                                                                                                       FEATURE:
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Gaps

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NAME/KEY: misc feature
LOCATION: (21) ... (21)
OTHER INFORMATION: n stands for thymidine
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NAME/KEY:
NAME/KEY:
NOTHER INFORMATION: 2'-deoxy-2'-fluoro
FEATURE:
NAME/KEY: misc_feature
LOCATION: (15)..(15)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
FEATURE:
NAME/KEY: misc_feature
LOCATION: (18)..(18)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
NAME/KEY: misc_feature
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NAME/KEY: misc_feature
LOCATION: (1)...(1)
OTHER_INFORMATION: 2'-deoxy-2'-fluoro
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NAME/KEY: misc_feature
LOCATION: (3)..(6)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19 CACCGTTCATTCTAGAGCA 1
                                                                                                                                                                                                                                                                                                                                     ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 163
US-10-923-330-575/c
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                                                                                                                                                                                                                                                                SEQ ID NO 574
LENGTH: 21
                                                                                                                                                                                                                                                                                                                   TYPE: RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE:
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Sequence 574, Application US/10923330

Sequence 574, Application US/10923330

Sequence 574, Application US/10923330

Sequence 574, Application No. US20050153916A1

Sequence 574, Application No. US20050153916A1

SEPECANT: Sirna Therapeutics, Inc.

APPLICANT: McSaviggen, James

APPLICANT: McSaviggen, James

APPLICANT: McSaviggen, James

APPLICANT: McSaviggen, James

APPLICANT: MAN Interference Mediated Inhibition of Telomerase Gene

TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)

TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)

FILE REFERENCE: 400/209 (MHB02-708-C)

CURRENT APPLICATION NUMBER: DCT/US03/04088

PRIOR FILING DATE: 2004-05-24

PRIOR FILING DATE: 2004-05-24

PRIOR FILING DATE: 2004-06-24

PRIOR FILING DATE: 2004-06-16

PRIOR FILING DATE: 2004-04-16

PRIOR FILING DATE: 2004-04-16

PRIOR FILING DATE: 2004-04-16

PRIOR FILING DATE: 2003-11-24

PRIOR FILING DATE: 2003-11-24
                                                                                                                                                                                                                                                FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region
FEATURE:
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PRIOR FILING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR FILING DATE: 2003-02-20
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 768
SOFTWARE: Patentin version 3.3
LENGTH: 21
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LOCATION: (20)...(20)
OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 19; DB 1; Length 21; Pred. No. 1e+02; 0; Mismatches 0; Indels
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100.0%; Pred. No. ...
... 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; LOCATION: (21)...(21)
; OTHER INFORMATION: n stands for thymidine
US-10-923-330-573
                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: 2'-deoxy-2'-fluoro FEATURE:
NAME/KEY: misc_feature
COCATION: (14)..(14)
                                                                                                                                                                                                                                                                                                                     NAME/KEY: misc feature
LOCATION: (2). .. (5)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ION: (14)..(14)
INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: misc feature
LOCATION: (17)...(17)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   148 CCACCGTTCATTCTAGAGC 166
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                                                                                                                                                                                                                    ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 100.0
Matches 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                             FEATURE: NAME/KEY: misc_feature LOCATION: (10)...(10)
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NAME/KEY: misc_feature
                                                                                                                                                                                                     TYPE: RNA
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FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region
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APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Beigelman, Jemes
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE REPERBENCE: 400/209 (MBHB02-708-C)
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT PILING DATE: 2004-08-20
PRIOR APPLICATION NUMBER: PCT/US03/04088
PRIOR PLING DATE: 2004-05-24
PRIOR FILING DATE: 2004-05-24
PRIOR FILING DATE: 2004-07-17
PRIOR PLING DATE: 2004-07-17
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PRIOR FILING DATE: 2003-11-23
PRIOR APPLICATION NUMBER: US 10/444,853
PRIOR FILING DATE: 2003-05-23
PRIOR PLING DATE: 2003-02-20
PRIOR PILING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR FILING DATE: 2003-02-20
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 768
SOFTWARE: Patentin version 3.3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ..
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NAME/KEY: misc_feature
LOCATION: (20)...(20)
OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 21;
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4.2%; Score 19; DB 1; Length 21;
Best Local Similarity 100.0%; Pred. No. 1e+02;
Matches 19; Conservative 0; Mismatches 0; Indels
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FEATURE:
NAME/KEY: misc_feature
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LOCATION: (21)..(21)
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Sequence 576, Application US/10923330

Sequence 576, Application US/10923330

Publication No. US20050153916A1

GENERAL INFORMATION:

APPLICANT: Sirna Therapeutics, Inc.

APPLICANT: McSwiggen, James

APPLICANT: Beigelman, Leonid

TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                  - See File Wrapper or PALM.
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LOCATION: (20) ...(20)
OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 19; DB 1; Length 21;
Pred. No. 1e+02;
0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4.2%; Scor.
100.0%; Pred. No. 100.0%; O; Mismatches
PRIOR FILING DATE: 2004-05-24
PRIOR APPLICATION NUMBER: US 10/826,966
PRIOR FILING DATE: 2004-04-16
PRIOR PELLON STATE 2004-01-16
PRIOR PELLON STATE 2004-01-14
PRIOR FILING DATE: 2004-01-14
PRIOR FILING DATE: 2003-11-23
PRIOR FILING DATE: 2003-11-23
PRIOR FILING DATE: 2003-11-23
PRIOR FILING DATE: 2003-05-23
PRIOR FILING DATE: 2003-05-23
PRIOR FILING DATE: 2003-05-23
PRIOR FILING DATE: 2003-05-20
PRIOR PILING DATE: 2003-05-20
PRIOR FILING DATE: 2003-05-20
PRIOR FILING DATE: 2003-02-20
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CTHER INFORMATION: n stands for thymidine US-10-923-330-575
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NAME/KEY: misc_feature
LOCATION: (1)..(2)
OTHER_INFORMATION: 2'-deoxy-2'-fluoro
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NAME/KEY: misc_feature
LOCATION: (4)..(7)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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NAME/KEX: misc_feature
LOCATION: (19)..(19)
OTHER_INFORMATION: 2'-deoxy-2'-fluoro
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NAME/KEX: misc_feature
LOCATION: (16)..(16)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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Best Local Similarity 100.
Matches 19, Conservative
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NAME/KEY: misc_feature
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OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region
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CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT PILING DATE: 2004-06-20
PRIOR APPLICATION NUMBER: US 60/396,600
PRIOR FILING DATE: 2004-05-24
PRIOR PILING DATE: 2004-05-24
PRIOR PILING DATE: 2004-05-24
PRIOR PILING DATE: 2004-04-16
PRIOR PILING DATE: 2004-04-16
PRIOR PILING DATE: 2004-04-16
PRIOR PILING DATE: 2004-04-16
PRIOR PILING DATE: 2003-11-24
PRIOR PILING DATE: 2003-11-23
PRIOR PILING DATE: 2003-12-23
PRIOR PILING DATE: 2003-05-23
PRIOR PILING DATE: 2003-05-23
PRIOR PILING DATE: 2003-02-20
PRIOR PILING DATE: 2003-03-20
PRIOR PILING DATE: 2003-03-20
PRIOR PILING DATE: 2003-03-20
PRIOR PILING DATE: 2003-03-20
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LOCATION: (20) ... (20)
OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
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4.2%; Score 19; DB 1; Les
Best Local Similarity 100.0%; Pred. No. 1e+02;
Matches 19; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ) OTHER INFORMATION: n stands for thymidine US-10-923-330-576
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: 2'-deoxy-2'-fluoro
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NAME/KEY: misc_feature
LOCATION: (4). . . (4)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LOCATION: (14) ... (19)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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Sequence 578, Application US/10923330

Publication No. US20050153916A1

GENERAL INFORMATION:

APPLICANT: Sirna Therapeutics, Inc.
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LOCATION: (20)...(20)
OTHER INFORMATION: n stands for thymidine
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ORGANISM: Artificial Sequence
       NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: misc_feature
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APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: REXPESSION Using Short Interfering RNA (sink)
FILE REPERENCE: 400/209 (MRHB02-708-C)
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT FILING DATE: 2004-08-20
PRIOR FILING DATE: 2004-05-24
PRIOR FILING DATE: 2004-05-24
PRIOR APPLICATION NUMBER: US 60/396,600
PRIOR APPLICATION NUMBER: US 60/396,600
PRIOR APPLICATION NUMBER: US 10/826,966
PRIOR APPLICATION NUMBER: US 10/757,803
PRIOR APPLICATION NUMBER: US 10/757,803
PRIOR PILING DATE: 2004-01-14
PRIOR FILING DATE: 2004-01-14
PRIOR FILING DATE: 2003-11-24
PRIOR FILING DATE: 2003-11-24
PRIOR PLILING DATE: 2003-01-123
PRIOR PLILING DATE: 2003-01-23
PRIOR PLILING DATE: 2003-05-23
PRIOR PLILING DATE: 2003-05-23
PRIOR FILING DATE: 2003-05-23
PRIOR FILING DATE: 2003-05-20
PRIOR PRI
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LOCATION: (1)...(1)
OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
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LOCATION: (1). (5)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (7)...(8)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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NAME/KEY: misc_feature
LOCATION: (14)..(17)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (10)..(12)
TYTER INFORMATION: 2'-deoxy-2'-fluoro
FEATURE:
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ORGANISM: Artificial Sequence
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LOCATION: (6). . (6)
OTHER INFORMATION: 2'-deoxy
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NAME/KEY: misc_feature
LOCATION: (9).7(9)
OTHER INFORMATION: 2'-deoxy
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LOCATION: (13)...(13)
OTHER INFORMATION: 2'-deoxy
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LOCATION: (18)...(19)
JTHER INFORMATION: 2'-deoxy
FEATURE:
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Sequence 579, Application US/10923330
| Sequence 579, Application US/10923330
| Publication No. US20050153916A1
| GENERAL INFORMATION:
| APPLICANT: Sitna Therapeutics, Inc.
| APPLICANT: MCSWAGEN, James
| APPLICANT: MCSWAGEN, James
| APPLICANT: MCSWAGEN, James
| APPLICANT: MCSWAGEN, James
| APPLICANT: Beigelman, Leonid
| TITLE OF INVENTION: Expression USing Short Interfering RNA (sinA)
| FILE REPERENCE: 400/209 (MRHG2-708-C)
| FILE REPERENCE: 400/209 (MRHG2-708-C)
| FILE REPLICATION NUMBER: US/10/23,330
| CURRENT FILING DATE: 2004-0.6-20
| PRIOR APPLICATION NUMBER: PCT/US03/0408
| PRIOR PLING DATE: 2004-0.5-24
| PRIOR APPLICATION NUMBER: PCT/US03/0408
| PRIOR PLING DATE: 2004-0.1-10
| PRIOR PLING DATE: 2004-0.1-12
| PRIOR PLING DATE: 2004-0.1-14
| PRIOR PLILING DATE: 2003-11-12
| PRIOR PLILING DATE: 2003-11-12
| PRIOR PLILING DATE: 2003-0.1-13
| PRIOR PLILING DATE: 2003-0.2-20
| PRIOR PLILING DATE: 2003-0.2-20
| PRIOR FILING DATE: 2003-0.2-20
| PRIOR PLILING DATE: 2003-0.2-20
| PRIOR FILING DATE: 2003-0.2-20
| PRIOR PRIO
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FEATURE:
NAME/KEY: misc_feature
LOCATION: (21)...(21)
CTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity
US-10-923-330-578
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LOCATION: (1)..(1)
OCHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
FEATURE:
NAME/KEY: misc feature
LOCATION: (1)..(4)..(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(2)...(
                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
4.2%; Score 19; DB 1; Length 21;
Best Local Similarity 63.2%; Pred. No. 1e+02;
Matches 12; Conservative 7; Mismatches 0; Indels
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OTHER INFORMATION: 2'-deoxy
FEATURE:
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LOCATION: (20)...(20)
OTHER INFORMATION: n stands for thymidine
                                                                                                                                                                                                                                                                                                                                                                                             LOCATION: (5). (6)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LOCATION: (8)..(10)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                NAME/KEY: misc_feature
LOCATION: (1)._(3)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      147 TCCACCGTTCATTCTAGAG 165
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                                                                                                          TYPE: RNA ORGANISM: Artificial Seguence
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Best Local Similarity 68.4%;
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                              NAME/KEY: misc_feature
LOCATION: (4).7(4)
OTHER INFORMATION: 2'-deoxy
                                                                                                                                                                                                                                                                                                                                                                                                                                                           LOCATION: (7). (7)
OTHER INFORMATION: 2'-deoxy
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OTHER INFORMATION: 2'-deoxy
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OTHER INFORMATION: 2'-deoxy
                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: misc_feature
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 4.2%; Score 19; DB 1; Length 21; Best Local Similarity 63.2%; Pred. No. 1e+02; Matches 12; Conservative 7; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                               FEATURE:
NAME/KEY: misc_feature
LOCATION: (20)...(20)
OTHER INFORMATION: n stands for thymidine
                                                                                                        FEATURE:
NAME/KEY: misc_feature
LOCATION: (9)..(11)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                  NAME/KEY: misc feature
LOCATION: (13) ... (16)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                LOCATION: (6).7(7)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      146 TTCCACCGTTCATTCTAGA 164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 UUCCACCGUUCAUUCUAGA 19
                                          FEATURE:
NAME/KEY: misc_feature
LOCATION: (8)...(8)
OTHER INFORMATION: 2'-deoxy
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OTHER INFORMATION: 2'-deoxy
                                                                                                                                                                                                                                                                                                                                LOCATION: (17)...(19)
OTHER INFORMATION: 2'-deoxy
NAME/KEY: misc feature
                                                                                                                                                                                    NAME/KEY: misc_feature
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PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR FILING DATE: 2003-02-20
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 76 NOS: 76 SEQ ID NOS SOFTWARE: Patentin version 3.3
SEQ ID NO S80
LENGTH: 21
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                                                                                                                                                                                                                                                                                                         NAME/KEY: misc feature
LOCATION: (1)..(1)
OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
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; OTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity
US-10-923-330-580
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Pred. No. 1e+02;
6; Mismatches
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LOCATION: (19)..(19)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                              NAME/KEY: misc_feature
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### APPLICANT: Beigelman, Leonid
### TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
### TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
### CINEMATION: Expression Using Short Interfering RNA (siNA)
### CINEMAT FILIAG DATE: 2004-08-20
### CURRENT FILIAG DATE: 2004-05-24
### PRIOR PELIAGION NUMBER: US 60/396,600
### RIOR PELIAGION NUMBER: US 10/826,966
### PRIOR PELIAGION NUMBER: US 10/826,966
### RIOR PELIAGION NUMBER: US 10/757,803
### RIOR PELIAGION NUMBER: US 10/757,803
### RIOR PELIAGION NUMBER: US 10/693,059
### RIOR PELIAGION NUMBER: US 10/693,059
### RIOR PELIAGION NUMBER: US 10/693,059
### RIOR PELIAGION NUMBER: US 10/444,853
### RIOR PELIAGION NUMBER: US 10/3-1-24
### RIOR PELIAGION NUMBER: US 10/3-20
### RIOR PELIAGION NUMBER: US 10/3-20
### RIOR PELIAGION NUMBER: US 10/3-20
### PRIOR PELIAGION NUMBER: US 10/3-20
### PRIOR PELIAGION NUMBER: PCT/US03/05346
### RIOR PELIAGION NUMBER: PCT/US03/05346
### RIOR PELIAGION NUMBER: PCT/US03/05028
### RIOR PELIAGION NUMBER: PCT/US03/050346
### RIOR PELIAGION NUMBER: PCT/US03/
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEX: misc feature
LOCATION: (1). (1)
OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
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NAME/KEY: misc_feature
LOCATION: (1)..(2)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (4)..(5)
OTHER_INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (7). (9)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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NAME/KEY: misc_feature
LOCATION: (11)..(14)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: RNA
ORGANISM: Artificial Sequence
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LOCATION: (3)...(3)
OTHER INFORMATION: 2'-deoxy
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LOCATION: (6). (6)
OTHER INFORMATION: 2'-deoxy
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OTHER INFORMATION: 2'-deoxy
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OTHER INFORMATION: 2'-deoxy
FEATURE:
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RESULT 16 Sequence 582, Application U8/10923330

Sequence 582, Application U8/10923330

Sequence 582, Application U8/10923330

Sequence 582, Application No. U920050153916A1

GENERAL INFORMATION: Largetuics, Inc.

APPLICANT: McSwiggen, James

APPLICANT: Legalman, Leonid

TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)

TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)

CURRENT APPLICATION WUMBER: DG/10/923, 330

FRIOR FILING DATE: 2004-05-24

PRIOR FILING DATE: 2004-05-24

PRIOR PELLOR TON WUMBER: DG/10/924, 639

PRIOR PELLOR DATE: 2004-05-24

PRIOR PELLOR DATE: 2004-06-24

PRIOR PELLOR DATE: 2004-01-14

PRIOR PELLOR DATE: 2003-01-23

PRIOR PELLOR DATE: 2003-01-23

PRIOR PELLOR DATE: 2003-01-23

PRIOR PELLOR DATE: 2003-02-20

PELOR PELLOR DATE: 2003-02-20

PELOR PELLOR DATE: 2003-02-20

PELOR PELOR PELOR DATE: 2003-02-20

PEROPERATION PELOR DATE: 2003-02-20

PEROPERATION PELON DATE: 2003-02-20

PELOR PELOR PELOR DATE: 2003-02-20

PEROPERATION PEROPERATION 
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                                                                                                                                                                                            NAME/KEY: misc_feature; LOCATION: (21)...(21); OTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity US-10-923-330-581
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NAME/K
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4.2%; Score 19; DB 1; Length 21;
Best Local Similarity 73.7%; Pred. No. 1e+02;
Matches 14; Conservative 5; Mismatches 0; Indels
LOCATION: (20)...(20)
OTHER INFORMATION: n stands for thymidine
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NAME/KEY: misc_feature
LOCATION: (1)...(1)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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ORGANISM: Artificial Sequence
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NAME/KEY: misc_feature
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4.2%; Score 19;
LING DATE: 2004-04-16
PLICATION NUMBER: US 10/757,803
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ORGANISM: Artificial Sequence
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OTHER INFORMATION: 2'-deoxy
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OTHER INFORMATION: 2'-deoxy
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OTHER INFORMATION: 2'-deoxy
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OTHER INFORMATION: 2'-deoxy
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NAME/KEY: misc_feature
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NAME/KEY: misc_feature
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LOCATION: (4)...(4)
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LOCATION: (5)...(7)
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NAME/KEY: misc_feature
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Publication No. US20050153916A1

GENERAL INFORMATION:
APPLICANT: Birna Therapeutics, Inc.
APPLICANT: Bedgelman, Leonid
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE REFERENCE: 400/209 (MBHB02-708-C)
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT FILING DATE: 2004-08-20
FRIOR APPLICATION NUMBER: PCT/US03/04088
FRIOR APPLICATION NUMBER: PCT/US03/04088
FRIOR APPLICATION NUMBER: CTT/US03/04088
FRIOR APPLICATION NUMBER: US 60/396,600
FRIOR FILING DATE: 2004-05-24
FRIOR APPLICATION NUMBER: US 10/8-24
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i LOCATION: (21)...(21)
i USCATION: 3'-3 attached terminal deoxyabasic moeity
US-10-923-330-582
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4.2%; Score 19; DB 1; Length 21;
Best Local Similarity 73.7%; Pred. No. 1e+02;
Matches 14; Conservative 5; Mismatches 0; Indele
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LOCATION: (20) ... (20)
OTHER INFORMATION: n stands for thymidine
                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: misc feature
LOCATION: (10)...(13)
JTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY: misc feature
LOCATION: (18) ... (18)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                    FEATURE:
NAME/KEY: misc_feature
LOCATION: (6)..(8)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                       LOCATION: (3). (4)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (2)..(2)

STHER INFORMATION: 2'-deoxy
                                                                                                                                                    NAME/KEY: misc_feature
LOCATION: (5)._(5)
OTHER INFORMATION: 2'-deoxy
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LOCATION: (14)...(17)
OTHER INFORMATION: 2'-deoxy
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THER INFORMATION: 2'-deoxy
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OTHER INFORMATION: 2'-deoxy
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OTHER INFORMATION: Description of Artificial Sequence: sinA sense region
FRIOR FILING DATE: 2004-01-14

FRIOR FILING DATE: 2004-01-124

FRIOR FILING DATE: 2003-11-24

FRIOR FILING DATE: 2003-11-24

FRIOR FILING DATE: 2003-11-23

FRIOR FILING DATE: 2003-11-23

FRIOR FILING DATE: 2003-05-23

FRIOR FILING DATE: 2003-05-23

FRIOR PELING DATE: 2003-05-20

FRIOR FILING DATE: 2003-05-20

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OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
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OTHER INFORMATION: n stands for thymidine
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (17)...(17)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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DB 1; Length 21;

US09436060A-16.rnpb.sl

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OTHER INFORMATION: 2'-deoxy
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LOCATION: (1).7(1)
OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
                                             0; Indels
73.7%; Pred. No. 1e+02;
tive 5; Mismatches
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LOCATION: (7)..(8)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (12)...(15)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                         150 ACCGTTCATTCTAGAGCAA 168
                                                                                                                                                   TYPE: RNA
ORGANISM: Artificial Sequence
   Best Local Similarity 73.74
Matches 14; Conservative
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NAME/KEY: misc feature
LOCATION: (1). (6)
OTHER INFORMATION: 2'-deoxy
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NAME/KEY: misc feature
LOCATION: (9)...(11)
OTHER INFORMATION: 2'-deoxy
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LOCATION: (16)..(16)
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LENGTH: 21
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## APPLICANY: Belgelann, Leoning
### APPLICANY: Belgelann, Leoning
### APPLICANY: Belgelann, Leoning
### TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
### FILE REPRENCE: 400/209 (MRHBRD2-708-C)
### CURRENT APPLICATION NUMBER: US/10/923,330
### CURRENT PILING DATE: 2004-08-20
### PRIOR PILING DATE: 2004-08-10
### PRIOR PILING DATE: 2004-08-10
### PRIOR PILING DATE: 2002-07-17
### PRIOR PILING DATE: 2004-08-16
### PRIOR PILING DATE: 2003-11-24
### PRIOR PILING DATE: 2003-02-20
### PRIOR PILING
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                                                                                                                                                                                                                                                                                                                            NAME/KEY: misc_feature

LOCATION: (20)

COTHER INFORMATION: n stands for thymidine

FEATURE:
NAME/KEY: misc_feature

LOCATION: (21)

COTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity
US-10-923-330-584
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 19; DB 1;
Pred. No. 1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5; Mismatches
NAME/KEY: misc feature
LOCATION: (17) ... (18)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 586, Application US/10923330
Publication No. US20050153916A1
GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
APPLICANT: Beigelman, Leonid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                300 GAAGAGTTGGGCTCTGTCA 318
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 73.7%;
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: RNA
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                        NAMES/KEY: misc feature
LOCATION: (19)...(19)
OTHER INFORMATION: 2'-deoxy
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OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region
               PRIOR FILING DATE: 2003-11-23
PRIOR APPLICATION NUMBER: US 10/444,853
PRIOR APPLICATION NUMBER: US 10/444,853
PRIOR FILING DATE: 2003-05-23
PRIOR FILING DATE: 2003-02-20
PRIOR FILING DATE: 2003-02-20
PRIOR FILING DATE: 2003-02-20
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 768
SOFTWARE: Patentin version 3.3
LENGTH: 21
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COATION: (20)...(20)

OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
FEATURE:
NAME/KEY: misc_feature

LOCATION: (21)...(21)

COTHER INFORMATION: n stands for thymidine

US-10-923-330-587
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APPLICATION NUMBER: US 10/693,059
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LOCATION: (8)..(8)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (12)...(12)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LOCATION: (1)..(3)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: misc feature
LOCATION: (9)..(11)
OTHER INFORMATION: 2'-deoxy
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OTHER INFORMATION: 2'-deoxy
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OTHER INFORMATION: 2'-deoxy
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LOCATION: (8). (8)
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LOCATION: (16)..(19)
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US-10-923-330-588/c
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                                                                                                                                                                                                                                                                                                                              TYPE: RNA
                                                                                                                                                                                                                                                                                                                                                                             FEATURE:
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Publication No. US20050153916A1
GENERAL INPORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
FILE REFERENCE: 400/209 (MBHB02-708-C)
CURRENT PAPLICATION NUMBER: US/10/923,330
CURRENT PILING DATE: 2004-08-20
PRIOR APPLICATION NUMBER: PCT/US03/0408
PRIOR PILING DATE: 2000-07-17
PRIOR PILING DATE: 2000-07-17
PRIOR PLING DATE: 2000-05-24
PRIOR PELING DATE: 2000-07-17
PRIOR PELING DATE: 2000-05-24
PRIOR PELING DATE: 2000-05-24
PRIOR PELING DATE: 2000-05-46
PRIOR PELING DATE: 2000-05-46
PRIOR PELING DATE: 2000-05-46
PRIOR PELING DATE: 2000-05-416
PRIOR FILING DATE: 2000-01-416
PRIOR FILING DATE: 2000-01-414
PRIOR FILING DATE: 2000-01-414
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NAME/KEY: misc_feature
LOCATION: (20) (20) (20)
OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4.2%; Score 19; DB 1; Length 21; 100.0%; Pred. No. 1e+02; tive 0; Mismatches 0; Indels
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// LOCATION: (21)...(21)

OTHER INFORMATION: n stands for thymidine

US-10-923-330-586
                                                                                                                        FEATURE:
NAME/KEY: misc_feature
LOCATION: (7)..(7)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                              NAME/KEY: misc feature
LOCATION: (11) ... (11)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: 2'-deoxy-2'-fluoro
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19 CTTCCACCGTTCATTCTAG 1
                                                                           LOCATION: (3). (6)
OTHER INFORMATION: 2'-deoxy
                                                                                                                                                                                                                                                NAME/KEY: misc_feature
LOCATION: (8)...(10)
OTHER INFORMATION: 2'-deoxy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LOCATION: (12)...(13)
OTHER INFORMATION: 2'-deoxy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LOCATION: (15)<sup>-</sup>.(19)
OTHER INFORMATION: 2'-deoxy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 100.0
Matches 19; Conservative
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NAME/KEY: misc_feature
                          FEATURE:
NAME/KEY: misc_feature
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Gaps

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OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region
                                                                                                                                                                   Length 21;
                                                                                                                                                              Query Match
4.2%; Score 19; DB 1; Length 21;
Best Local Similarity 100.0%; Pred. No. 1e+02;
Matches 19; Conservative 0; Mismatches 0; Indels
       ; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (21)...(21)
. CTHER INFORMATION: n stands for thymidine
US-10-923-330-588
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LOCATION: (2)...(5)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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NAME/KEY: misc feature
LOCATION: (10) (10)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                             147 TCCACCGTTCATTCTAGAG 165
                                                                                                                                                                                                                                                                                            19 TCCACCGTTCATTCTAGAG 1
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ORGANISM: Artificial Sequence
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LOCATION: (1). (1)
OTHER INFORMATION: 2'-deoxy
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OTHER INFORMATION: 2'-deoxy
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NAME/KEY: misc_feature
LOCATION: (6). . (9)
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          APPLICANT: Brancapeurics, Inc.
APPLICANT: McSwiggen, James
APPLICANT: Beigelman, Leonid
ITILE OF INVENTION: Experession Using Short Interfering RNA (sinA)
FILE BRENERNCE: 400/209 (WHBNO-709-C)
FILE OF INVENTION: Expression Using Short Interfering RNA (sinA)
FILE REPERENCE: 400/209 (WHBNO-709-C)
CURRENT APPLICATION UNMER: US/10/923,330
CURRENT FILING DATE: 2004-06-24
PRIOR APPLICATION NUMBER: US 60/396,600
PRIOR PELING DATE: 2004-05-24
PRIOR PLING DATE: 2004-06-24
PRIOR PLING DATE: 2004-06-14
PRIOR PLING DATE: 2004-06-14
PRIOR PLING DATE: 2004-06-14
PRIOR PLING DATE: 2003-01-14
PRIOR PLING DATE: 2003-01-23
PRIOR PLING DATE: 2003-02-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region
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LOCATION: (20)...(20)
OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)._(4)
OTHER_INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
NAME/KEY: misc_feature
LOCATION: (9)..(9)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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NAME/KEY: misc_feature
LOCATION: (13)...(13)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (16)...(16)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
APPLICANT: Sirna Therapeutics, Inc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: RNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: misc feature
LOCATION: (5). (8)
OTHER INFORMATION: 2'-deoxy
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LOCATION: (10)...(12)
OTHER INFORMATION: 2'-deoxy
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LOCATION: (14).(15)
OTHER INFORMATION: 2'-deoxy
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NAME/KEY: misc_feature
LOCATION: (17)...(19)
OTHER INFORMATION: 2'-deoxy
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OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FEATURE:

NAME/KEX: misc_feature
LOCATION: (15)-.(15)

OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (18)...(18)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (1)._(1)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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NUMBER OF SEQ ID NOS: 768
SOFTWARE: Patentin version 3.3
SEQ ID NO 590
LENGTH: 21
                                                                                                                           ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                              NAME/KEY: misc_feature
LOCATION: (2)._(2)
OTHER INFORMATION: 2'-deoxy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LOCATION: (7). (10)
OTHER INFORMATION: 2'-deoxy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LOCATION: (12)...(14)
OTHER INFORMATION: 2'-deoxy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LOCATION: (16)...(17)
OTHER INFORMATION: 2'-deoxy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LOCATION: (19)...(19)
OTHER INFORMATION: 2'-deoxy
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LOCATION: (11)..(11)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         LOCATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE:
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US-10-231-330-590/C

US-10-231-330-590/C

Publication No. US20050153916A1

GENERAL INFORMATION:

APPLICANT: Sirna Therapeutics, Inc.

APPLICANT: McSwiggen, James

APPLICANT: McSwiggen, James

TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene

TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene

TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene

TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene

TITLE OF INVENTION: RNA INTERFERENCE: 2004-08-20

CURRENT FILING DATE: 2004-08-20

FRIOR FILING DATE: 2004-06-24

FRIOR FILING DATE: 2004-05-24

FRIOR PILING DATE: 2004-05-24

FRIOR FILING DATE: 2004-01-14

FRIOR FILING DATE: 2004-01-14

FRIOR FILING DATE: 2004-01-14

FRIOR FILING DATE: 2003-11-23

FRIOR FILING DATE: 2003-11-23

FRIOR FILING DATE: 2003-11-23

FRIOR FILING DATE: 2003-11-23

FRIOR FILING DATE: 2003-02-13

FRIOR FILING DATE: 2003-02-23

FRIOR FILING DATE: 2003-02-20

FRIOR FILING DATE: 2003-02-20
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NAME/KEY: misc feature
LOCATION: (18)...(19)
OTHER INFORMATION: 2'-deoxy
PRATURE:
NAME/KEY: misc feature
LOCATION: (20)...(20)
OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 4.2%; Score 19; DB 1; Length 21; Best Local Similarity 100.0%; Pred. No. 1e+02; Matches 19; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: misc_feature

// LOCATION: (21)...(21)

// OTHER INFORMATION: n stands for thymidine

US-10-923-330-589
                                                                                                                                                                                                                                                                                                                           NAME/KEY: misc feature
LOCATION: (17) ... (17)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                   FEATURE:
NAME/KEX: misc_feature
LOCATION: (14)..(14)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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                                                         LOCATION: (11)..(13)
OTHER INFORMATION: 2'-deoxy
                                                                                                                                                                                                                          NAME/KEY: misc_feature
LOCATION: (15)...(16)
OTHER INFORMATION: 2'-deoxy
FEATURE:
FEATURE: MAME/KEY: misc feature LOCATION: (11)...(13)
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LOCATION: (20) .. (20) OTHER INFORMATION: Phosphorothicate 3'-Internuclectide Linkage
                                                                                                                                                                                                                  Query Match 4.2%; Score 19; DB 1; Length 21; Best Local Similarity 100.0%; Pred. No. 1e+02; Matches 19; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-10-923-330-591/c
; Sequence 591, Application US/10923330
; Publication No. US20050153916A1
; GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
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NAME/KEY: misc_feature
LOCATION: (20)...(20)
OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
FEATURE:
NAME/KEY: misc_feature
LOCATION: (21)...(21)
OTHER INFORMATION: n stands for thymidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 592, Application US/10923330 Publication No. US20050153916Al GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                      150 ACCGTTCATTCTAGAGCAA 168
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
NAME/KEY: misc_feature
LOCATION: (2)...(3)
OTHER INFORMATION: 2'-deoxy
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       APPLICANT: MCSALGABL, Jeonid
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE REPERENCE: 400/209 (WHHB02-708-C)
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT APPLICATION NUMBER: US 60/396,600
PRIOR FILING DATE: 2004-05-24
PRIOR PILING DATE: 2004-05-24
PRIOR PELING DATE: 2004-05-24
PRIOR PELING DATE: 2004-05-24
PRIOR PELING DATE: 2004-05-24
PRIOR PLING DATE: 2004-05-34
PRIOR PLING DATE: 2004-01-16
PRIOR PELING DATE: 2004-01-14
PRIOR PLING DATE: 2004-01-14
PRIOR PLING DATE: 2004-01-14
PRIOR PLING DATE: 2004-01-12
PRIOR PLING DATE: 2003-01-12
PRIOR PLING DATE: 2003-01-12
PRIOR PLING DATE: 2003-01-12
PRIOR PLING DATE: 2003-01-23
PRIOR PLING DATE: 2003-02-23
PRIOR PLING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: US 10/693,059
PRIOR PLING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: US 10/693,059
PRIOR PLING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: US 10/693,059
PRIOR PLING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: PCT/US03/05346
PRIOR PLING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: PCT/US03/05346
PRIOR PLING DATE: 2003-02-20
PRIOR PLING DATE: 2003-03-33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE: OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: misc feature
LOCATION: (1)...(2)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (4). 7)
JTHER INFORMATION: 2'-deoxy-2'-fluoro
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NAME/KEY: misc_feature
LOCATION: (12)..(12)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (16)...(16)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (19)..(19)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: RNA
ORGANISM: Artificial Sequence
McSwiggen, James
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LOCATION: (3)._(3)
OTHER INFORMATION: 2'-deoxy
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LOCATION: (13)...(15)
JTHER INFORMATION: 2'-deoxy
FEATURE:
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LOCATION: (8)...(11)
OTHER INFORMATION: 2'-deoxy
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NAME/KEY: misc_feature
LOCATION: (17)..(18)
OTHER INFORMATION: 2'-deoxy
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APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, James

TITLE 0F INVENTION: RAP Interference Mediated Inhibition of Telomerase Gene
TITLE 0F INVENTION: RAPPLICANTON WINBER: US,10/923,330
CURRENT APPLICATION WINBER: US,10/923,330
CURRENT APPLICATION WINBER: US,204-05-24
PRIOR PLING DATE: 2004-05-24
PRIOR PLING DATE: 2004-05-24
PRIOR PLING DATE: 2004-05-24
PRIOR APPLICATION WINBER: US 10/826,966
PRIOR PLING DATE: 2004-06-16
PRIOR APPLICATION WINBER: US 10/757,803
PRIOR PLING DATE: 2004-01-14
PRIOR PLING DATE: 2004-01-14
PRIOR PLING DATE: 2004-01-14
PRIOR PLING DATE: 2003-01-23
PRIOR PLING DATE: 2003-01-23
PRIOR PLING DATE: 2003-01-23
PRIOR PLING DATE: 2003-01-23
PRIOR PLING DATE: 2003-02-20
PRIOR APPLICATION WINBER: US 10/753,659
PRIOR PLING DATE: 2003-02-20
PLING DATE: 2003-02
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NAME/KEY: misc_feature
LOCATION: (1) .. (1) .. (1) OTHER INFORMATION: 2'-deoxy-2'-fluoro
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Query Match 4.2%; Score 19; DB 1; Length 21; Best Local Similarity 100.0%; Pred. No. 1e+02; Matches 19; Conservative 0; Mismatches 0; Indels
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
FEATURE:
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ORGANISM: Artificial Sequence
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NAME/KEY: misc_feature
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LOCATION: (14)..(17)
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RESULT 170

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LOCATION: (20) ...(20)
OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 4.2%; Score 19; DB 1; Length 21; Best Local Similarity 100.0%; Pred. No. 1e+02; Matches 19; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FEATURE:

NAME/KEY: misc_feature

LOCATION: (21) - (21)

OTHER INFORMATION: n stands for thymidine
US-10-923-330-592
                                                                                                                            NAME/KEY: misc_feature
LOCATION: (9)._(11)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                   LOCATION: (14)...(19)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GAAGAGTTGGGCTCTGTCA 1
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NAME/KEY: misc_feature
LOCATION: (12)...(13)
OCTHER INFORMATION: 2'-deoxy
FEATURE:
NAME/KEY: misc_feature
LOCATION: (5)._(8)
OTHER INFORMATION: 2'-deoxy
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LENGTH: 21
TYPE: RNA
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Sequence 595, Application US/10923330
Publication No. US20050153916A1
GENERAL INFORMATION:
BELL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE REFERENCE: 400/209 (MBHB02-708-C)
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT FILING DATE: 2004-08-20
PRIOR FILING DATE: 2004-05-24
OTHER INFORMATION: Description of Artificial Sequence: siNA sense region
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CTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity
US-10-923-330-594
                                                                                      LOCATION: (1). (1) (1) OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 4.2%; Score 19; DB 1; Length 21; Best Local Similarity 63.2%; Pred. No. 1e+02; Matches 12; Conservative 7; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: n stands for thymidine
                                                                                                                                                                                                                                                                                                                                                                                                                   NAME/KEY: misc feature
LOCATION: (7). [8)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: misc feature
LOCATION: (10) ... (12)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                           LOCATION: (1). (5)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LOCATION: (14)..(17)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   145 CTTCCACCGTTCATTCTAG 163
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1 CUUCCACCGUUCAUUCUAG 19
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LOCATION: (9)._(9)
OTHER INFORMATION: 2'-O-methyl
                                                                                                                                                                                                                                                                                                                                LOCATION: (6). T(6)
OTHER INFORMATION: 2'-0-methyl
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OTHER INFORMATION: 2'-0-methyl
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OTHER INFORMATION: 2'-0-methyl
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WESULT 18

WESULT 18

WESULT 18

WESULT 18

Sequence 596, Application US/10923330

Sequence 596, Application US/10923330

Sequence 596, Application No. US20050153916A1

APPLICANT: Sinta Therapeutics, Inc.

APPLICANT: Hostagelman, Leonid

TITLE OF INVENTION: Expression Using Short Interfering RNA (SiNA)

TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene

TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene

FILE REFRENCE: 400/209 (MEHH02/209.)

CURRENT APLICATION NUMBER: DC10/408.20

PRIOR APLICATION NUMBER: US 10/408.20

PRIOR APLICATION NUMBER: US 10/40.40

PRIOR PELING DATE: 2004-00-24

PRIOR PELING DATE: 2004-00-14

PRIOR PELING DATE: 2004-00-11-24

PRIOR PELING DATE: 2004-01-14

PRIOR PELING DATE: 2004-01-14

PRIOR PILING DATE: 2004-01-14

PRIOR PELING DATE: 2004-01-14

PRIOR PELING DATE: 2003-11-23

PRIOR PELING DATE: 2003-11-23

PRIOR PELING DATE: 2003-01-23

PRIOR PELING DATE: 2003-01-23

PRIOR PELING DATE: 2003-01-23

PRIOR PELING DATE: 2003-01-23

PRIOR APPLICATION NUMBER: US 10/693,059

PRIOR PELING DATE: 2003-02-20

PRIOR PELING DATE: 2003-
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                                                                                                     Gaps
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OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
Query Match 4.2%; Score 19; DB 1; Length 21; Best Local Similarity 63.2%; Pred. No. 1e+02; Matches 12; Conservative 7; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)..(3)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: misc feature
LOCATION: (5).7(6)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                    146 TTCCACCGTTCATTCTAGA 164
                                                                                                                                                                                                       UCCACCGUUCAUUCUAGA 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LOCATION: (4)...(4)
OTHER INFORMATION: 2'-0-methyl
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY: misc_feature
LOCATION: (7)..(7)
OTHER INFORMATION: 2'-O-methyl
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: RNA
ORGANISM: Artificial Sequence
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NAME/KEY: misc_feature
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NAME/KEY: misc_feature
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APPLICANT: MCSWiggen, James
APPLICANT: McSwiggen, James
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: RAM Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE REFERENCE: 400/209 (MRHSD2-708-C)
CURRENT APPLICATION NUMBER: US/10/923,330
                                   siNA sense region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ô
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                                                                                                                           LOCATION: (1)...(1)
OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
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FEATURE:
OTHER INFORMATION: Description of Artificial Sequence:
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4.2%; Score 19; DB 1;
Best Local Similarity 73.7%; Pred. No. 1e+02;
Matches 14; Conservative 5; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: misc feature
LOCATION: (20) ... (20)
OTHER INFORMATION: n stands for thymidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LOCATION: (4). (5)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LOCATION: (11)...(14)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                   NAME/KEY: misc feature
LOCATION: (1)._(2)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LOCATION: (19)...(19)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 598, Application US/10923330
Publication No. US20050153916A1
GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             148 CCACCGTTCATTCTAGAGC 166
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OTHER INFORMATION: 2'-0-methyl
                                                                                                                                                                                                                                                                                                                                                                                   LOCATION: (3).7(3)

OTHER INFORMATION: 2'-O-methyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LOCATION: (6)..(6)
OTHER INFORMATION: 2'-0-methyl
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OTHER INFORMATION: 2'-O-methyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PEATURE:
NAME/KEY: misc_feature
                                                                                              NAME/KEY: misc_feature
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LOCATION: (6)..(6)
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LOCATION: (7)..(9)
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APPLICANT: McSwiggen, James
APPLICANT: Beigelman, Leonid

TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE REFERENCE: 400/209 (WBBNB2-708-C)
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT APPLICATION NUMBER: US/04-08-20
PRIOR APPLICATION NUMBER: US 60/396,600
PRIOR PLING DATE: 2004-05-24
PRIOR PLING DATE: 2004-05-24
PRIOR PLING DATE: 2004-05-24
PRIOR PLING DATE: 2004-06-10
PRIOR PLING DATE: 2004-06-10
PRIOR PLING DATE: 2004-06-10
PRIOR PLING DATE: 2004-06-114
PRIOR PLING DATE: 2004-06-114
PRIOR PLING DATE: 2004-01-14
PRIOR PLING DATE: 2003-11-24
PRIOR PLING DATE: 2003-11-24
PRIOR PLING DATE: 2003-11-23
PRIOR PLING DATE: 2003-10-14
PRIOR PLING DATE: 2003-01-22
PRIOR PLING DATE: 2003-02-20
PRIOR PLING PLING DATE: 2003-02-20
PRIOR PLING 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: misc_feature

1 LOCATION: (21)...(21)

2 OTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity

US-10-923-330-596
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 19; DB 1; Length 21;
Pred. No. 1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                6; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LOCATION: (20)...(20)
OTHER INFORMATION: n stands for thymidine
                                                                                                                                                                                      FEATURE:
NAME/KEY: misc_feature
LOCATION: (12) ... (15)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
   LOCATION: (8)..(10)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 597, Application US/10923330 Publication No. US20050153916A1 GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           147 TCCACCGTTCATTCTAGAG 165
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CCACCGUUCAUUCUAGAG 19
                                                                                                                                                                                                                                                                                                                  FEATURE:
NAME/KEX: misc_feature
LOCATION: (16)..(19)
OTHER INFORMATION: 2'-0-methyl
                                                                                                                              LOCATION: (11)...(11)
OTHER INFORMATION: 2'-O-methyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 68.4%;
Matches 13; Conservative
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ORGANISM: Artificial Sequence
                                                                                              NAME/KEY: misc feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: misc_feature
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Gaps

Page 78

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OTHER INFORMATION: Description of Artificial Sequence: sink sense region
PRIOR APPLICATION NUMBER: PCT/US03/04088

PRIOR PILING DATE: 2004-08-20

PRIOR FILING DATE: 2004-05-24

PRIOR PELLON NUMBER: US 60/396,600

PRIOR PELLON NUMBER: US 60/396,600

PRIOR PELLON NUMBER: US 10/826,966

PRIOR APPLICATION NUMBER: US 10/826,966

PRIOR APPLICATION NUMBER: US 10/757,803

PRIOR PILING DATE: 2004-01-14

PRIOR PILING DATE: 2004-01-14

PRIOR PILING DATE: 2004-01-14

PRIOR PILING DATE: 2003-11-24

PRIOR PELLON NUMBER: US 10/693,059

PRIOR PILING DATE: 2003-11-24

PRIOR PILING DATE: 2003-11-24

PRIOR PILING DATE: 2003-11-23

PRIOR PILING DATE: 2003-05-23

PRIOR PILING DATE: 2003-05-20

PRIOR APPLICATION NUMBER: PCT/US03/0508

PRIOR PILING DATE: 2003-05-23

PRIOR PILING DATE: 2003-05-23

PRIOR APPLICATION NUMBER: PCT/US03/0508

PRIOR PILING DATE: 2003-05-23

PRIOR PILING DATE: 2003-05-30

PRIOR PILING DATE: 200
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LOCATION: (1). (1)
OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
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LOCATION: (1).7(1)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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NAME/KEY: misc_feature
LOCATION: (18) .. (18)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER_INFORMATION: 2'-deoxy-2'-fluoro
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THER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (2)...(2)
OTHER INFORMATION: 2'-0-methyl
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LOCATION: (9)...(9)
OTHER INFORMATION: 2'-O-methyl
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ORGANISM: Artificial Sequence
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LOCATION: (5)._(5)
OTHER INFORMATION: 2'-0-methyl
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LOCATION: (14)...(17)
OTHER INFORMATION: 2'-O-methyl
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LOCATION: (19)..(19)
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NESOUR. 18. Sequence 599, Application US/10923330

Sequence 599, Application US/10923330

Publication No. US20050153916A1

GENERAL INFORMATION:

APPLICANT: BESIGNATION:

APPLICANT: BESIGNATION:

APPLICANT: BESIGNATION: Expression Using Short Interfering RNA (sinA)

TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)

TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)

FILE REFERENCE: 400/209, (WHHEND2.709-C)

CURRENT APPLICATION NUMBER: US/10/923,330

PRIOR APPLICATION NUMBER: US/10/924,088

PRIOR APPLICATION NUMBER: US/10/924,088

PRIOR PELING DATE: 2004-05-24

PRIOR APPLICATION NUMBER: US/10/926,600

PRIOR APPLICATION NUMBER: US/10/926,966

PRIOR APPLICATION NUMBER: US/10/926,966

PRIOR PELING DATE: 2004-04-16

PRIOR PELING DATE: 2004-04-16

PRIOR PELING DATE: US/10/92,966

PRIOR PELING DATE: US/10/92,966

PRIOR PELING DATE: US/10/92,966

PRIOR PELING DATE: US/10/92,966

PRIOR PELING DATE: 2003-11-23

PRIOR PELING DATE: 2003-11-24

PRIOR PELING DATE: 2003-11-23

PRIOR PELING DATE: 2003-11-24

PRIOR PELING DATE: 2003-05-23

PRIOR PELING DA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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0
                                                                    NAME/KEY: misc feature
LOCATION: (20) (20)
OTHER INFORMATION: n stands for thymidine
FEATURE:
NAME/KEY: misc feature
LOCATION: (21) (21)
OTHER INFORMATION: 3 - 3 attached terminal deoxyabasic moeity
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LOCATION: (1)..[1)
OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
FRATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
4.2%; Score 19; DB 1; Length 21;
Best Local Similarity 73.7%; Pred. No. 1e+02;
Matches 14; Conservative 5; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                149 CACCGTTCATTCTAGAGCA 167
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: 2'-0-methyl
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OTHER INFORMATION: 2'-0-methyl
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ORGANISM: Artificial Sequence
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NAME/KEY: misc_feature
LOCATION: (2)...(3)
                                                                                                                                                                                                                                                                                                                                                    US-10-923-330-598
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siNA sense region
               PRIOR FILING DATE: 2003-11-23
PRIOR FILING DATE: 2003-05-23
PRIOR FILING DATE: 2003-05-23
PRIOR FILING DATE: 2003-05-23
PRIOR FILING DATE: 2003-02-20
PRIOR FILING DATE: 2003-02-20
PRIOR APPLICATION WUMBER: PCT/US03/05028
PRIOR FILING DATE: 2003-02-20
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 768
SOFTWARE: Patentin version 3.3
SEQ ID NO 600
LENGTH: 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: misc_feature

LOCATION: (21)...(21)

CTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity

US-10-923-330-600
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LOCATION: (1)._(1)
OTHER INFORMATION: 5'-3 attached terminal deoxyabasic mosity
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                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: Description of Artificial Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4.2%; Score 19; DB 1; 73.7%; Pred. No. 1e+02; tive 5; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: n stands for thymidine
APPLICATION NUMBER: US 10/693,059
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LOCATION: (17)...(18)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LOCATION: (7). (8)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LOCATION: (12)..(15)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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US-10-923-330-602/c
US-10-923-330-602, Application US/10923330
; Publication No. US2050153916A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         300 GAAGAGTTGGGCTCTGTCA 318
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OTHER INFORMATION: 2'-0-methyl
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OTHER INFORMATION: 2'-0-methyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LOCATION: (16)...(16)
OTHER INFORMATION: 2'-0-methyl
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OTHER INFORMATION: 2'-O-methyl
                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 73.73
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: misc feature
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NAME/KEY: misc_feature
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NAME/KEY: misc_feature
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LOCATION: (20)...(20)
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                                                                                                                                                                                                                                                                                                                            TYPE: RNA
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sequence 600, Application US/10923330

publication No. US20050153916A1

GENERAL INFORMATION:

APPLICANT: Siran Therapeutics, Inc.

APPLICANT: Beigelman, Leonid

; TITLE OF INVENTION: RAN Interference Mediated Inhibition of Telomerase Gene

; TITLE OF INVENTION: RAN Interference Mediated Inhibition of Telomerase Gene

; TITLE OF INVENTION: RANDERS: US/10/923,330

; TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)

; FILE REFERENCE: 400/209 (MBHB02-708-C)

; FURRENT APPLICATION NUMBER: DCT/US03/04088

; PRIOR FILING DATE: 2004-05-24

; PRIOR PLING DATE: 2002-07-17

; PRIOR PLING DATE: 2004-06-24

; PRIOR FILING DATE: 2004-06-24

; PRIOR FILING DATE: 2004-06-24

; PRIOR FILING DATE: 2004-01-14

; PRIOR FILING DATE: 2003-11-24
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; LOCATION: (21)...(21)
; OTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity
US-10-923-330-599
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
4.2%; Score 19; DB 1; Length 21;
Best Local Similarity 73.7%; Pred. No. 1e+02;
Matches 14; Conservative 5; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FEATURE:
NAME/KEY: misc_feature
LOCATION: (20)..(20)
OTHER INFORMATION: n stands for thymidine
                                                                                                                    FEATURE:
NAME/KEY: misc_feature
LOCATION: (5)..(7)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                            NAMEXEY: misc feature
LOCATION: (9). (12)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY: misc feature
LOCATION: (17)...(17)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (18)..(19)
OTHER INFORMATION: 2'-O-methyl
                                              NAME/KEY: misc_feature
LOCATION: (4)._(4)
OTHER INFORMATION: 2'-0-methyl
                                                                                                                                                                                                                                            NAME/KEY: misc feature
LOCATION: (8). (8)
OTHER INFORMATION: 2'-0-methyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LOCATION: (13)..(16)
OTHER INFORMATION: 2'-0-methyl
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NAME/KEY: misc feature
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Gape

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; NAME/KEY: misc_feature
; LOCATION: (21)...(21)
; OTHER INFORMATION: n stands for thymidine
US-10-923-330-602
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NAME/KRY: misc_feature
LOCATION: (1) ...(3)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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                                                                                                                                                                                                                                                                                                                                                                                  19 CTTCCACCGTTCATTCTAG 1
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LOCATION: (4). (7)
OTHER INFORMATION: 2'-O-methyl
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ORGANISM: Artificial Sequence
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APPLICANT: SITUR INCEADEULICS, INC.
APPLICANT: McGwiggen, James
APPLICANT: Beigelman, Leonis
APPLICANT: Beigelman, Leonis
APPLICANT: Beigelman, Leonis
APPLICANT: Beigelman, Leonis
TITLE OF INVENTION: RAY INTEFFERENCE Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
FILE REFERENCE: 400/209 (MBHB02-708-C)
CURRENT APPLICATION NUMBER: 108/10/923,330
CURRENT FILING DATE: 2004-08-20
PRIOR APPLICATION NUMBER: PCT/US03/0408
PRIOR APPLICATION NUMBER: US 60/396,600
PRIOR FILING DATE: 2004-01-14
PRIOR PELING DATE: 2004-01-14
PRIOR PELING DATE: 2004-01-14
PRIOR PELING DATE: 2003-01-14
PRIOR PELING DATE: 2003-01-14
PRIOR PELING DATE: 2003-01-14
PRIOR APPLICATION NUMBER: US 10/757,803
PRIOR PELING DATE: 2003-11-24
PRIOR PELING DATE: 2003-11-24
PRIOR PELING DATE: 2003-11-24
PRIOR PELING DATE: 2003-05-23
PRIOR PELING DATE: 2003-05-23
PRIOR APPLICATION NUMBER: US 10/444,853
PRIOR PELING DATE: 2003-05-20
PRIOR PELING DATE: 2003-05-20
PRIOR PELING DATE: 2003-02-20
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LOCATION: (20)...(20)
OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: misc_feature
LOCATION: (1)...(2)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
FEATURE:
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (11)...(11)
JTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (14)...(14)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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NAME/KEY: misc_feature
LOCATION: (12)..(13)
OTHER INFORMATION: 2'-0-methyl
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OTHER INFORMATION: 2'-0-methyl
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NAME/KEY: misc_feature
LOCATION: (8)..(10)
OTHER_INFORMATION: 2'-0-methyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: RNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY: misc_feature
LOCATION: (15)..(19)
OTHER_INFORMATION: 2'-0-methyl
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LENGTH: 21
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| Publication No. U820050153916A1
| Publication U820050153916A1
| Publication No. U820050153916A1
| Publication No. U820050153916A1
| APPLICANT: Sirna Therapeutics, Inc.
| APPLICANT: Girna Therapeutics, Inc.
| APPLICANT: Beigelman, Leonid
| TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
| TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
| TITLE OF INVENTION: RNA INTERFERENCE: 400/209 (MRHE02-708-C)
| CURRENT PEPLICATION NUMBER: U8/10/933,330
| CURRENT FILING DATE: 2004-05-24
| PRIOR PAPLICATION NUMBER: DCT/U803/0408
| PRIOR PELING DATE: 2004-05-24
| PRIOR PAPLICATION NUMBER: US 60/396,600
| PRIOR PELING DATE: 2004-05-24
| PRIOR PAPLICATION NUMBER: US 10/720,448
| PRIOR PELING DATE: 2004-01-14
| PRIOR PELING DATE: 2004-01-12
| PRIOR PELING DATE: 2004-01-14
| PRIOR PELING DATE: 2003-01-12
| PRIOR PELING DATE: 2003-01-12
| PRIOR PELING DATE: 2003-01-12
| PRIOR PELING DATE: 2003-01-14
| PRIOR PELING DATE: 2003-01-13
| PRIOR PELING DATE: 2003-01-24
| PRIOR PELING DATE: 2003-01-34
| PRIOR PELING DATE: 2003-01-34
| PRIOR PELING DATE: 2003-01-34
| PRIOR PELING DATE: 2003-02-20
| PRIOR 
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Query Match
4.2%; Score 19; DB 1; Length 21;
Best Local Similarity 100.0%; Pred. No. 1e+02;
Matches 19; Conservative 0; Mismatches 0; Indels
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LOCATION: (9)...(11)
OTHER INFORMATION: 2'-0-methyl
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OTHER INFORMATION: Description of Artificial Sequence: sink antisense region
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APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Bigleman, Leon, James
APPLICANT: Beigelman, Leon, Therefore Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RNA Interference Mediated Interfering RNA (sina)
CURRENT APPLICATION NUMBER: US/10/923,330
PRIOR APPLICATION NUMBER: PCT/US03/0408
PRIOR APPLICATION NUMBER: US 60/396,600
PRIOR APPLICATION NUMBER: PCT/US04/16390
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NAME/KEY: misc_feature
LOCATION: (20) ... (20)
OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 4.2%; Score 19; DB 1; Length 21; Best Local Similarity 100.0%; Pred. No. 1e+02; Matches 19; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; LOCATION: (21) ... (21)
; OTHER INFORMATION: n stands for thymidine US-10-923-330-604
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME/KEY: misc feature
LOCATION: (13) ... (13)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: misc_feature
LOCATION: (9)._(9)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                    LOCATION: (1). (4)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LOCATION: (16)...(16)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 605, Application US/10923330 Publication No. US20050153916A1 GENERAL INFORMATION:
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NAME/KEY: misc feature
LOCATION: (10)...(12)
OTHER INFORMATION: 2'-O-methyl
FEATURE:
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LOCATION: (14)..(15)
OTHER INFORMATION: 2'-0-methyl
                                                                                                                                                                                                                                                                                                         LOCATION: (5).7(8)
OTHER INFORMATION: 2'-0-methyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LOCATION: (17)..(19)
OTHER INFORMATION: 2'-0-methyl
   ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                  NAME/KEY: misc_feature
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LOCATION: (16)...(16)
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NAME/KEY: misc_feature
                                                                                                                                 NAME/KEY: misc_feature
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GENERAL INCRAMINED

GENERAL INCRAMINED

APPLICANT: SITUR Therapeutice, Inc.

APPLICANT: McSwiggen, James

TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)

FILE REFRENCE: 400/209 (MHHD0-C)

CURRENT APPLICATION NUMBER: US/10/923,330

CURRENT FILING DATE: 2004-06-20

FRIOR FILING DATE: 2004-06-20

FRIOR FILING DATE: 2004-06-24

FRIOR FILING DATE: 2004-06-24

FRIOR FILING DATE: 2004-06-34

FRIOR FILING DATE: 2004-04-16

FRIOR FILING DATE: 2004-04-16

FRIOR FILING DATE: 2004-04-16

FRIOR FILING DATE: 2003-11-3

FRIOR FILING DATE: 2003-10-3

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LOCATION: (20) ...(20)
OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 19; DB 1; Length 21;
Pred. No. 1e+02;
0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 4.2%; Score 19; DB Best Local Similarity 100.0%; Pred. No. 1e+ Matches 19; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 : LOCATION: (21)...(21)
; OTHER INFORMATION: n stands for thymidine US-10-923-330-603
   NAME/KEY: misc_feature
LOCATION: (12) \(\tilde{\chi}\).(12)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                   FEATURE:
NAME/KEX: misc feature
LOCATION: (15)-(15)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-10-923-330-604/c; Sequence 604, Application US/10923330; Publication No. US20050153916A1; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TTCCACCGTTCATTCTAGA 1
                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: misc_feature
LOCATION: (16)..(19)
OTHER INFORMATION: 2'-O-methyl
                                                                                                 FEATURE:
NAME/KEX: misc_feature
LOCATION: (13)..(14)
OTHER_INFORMATION: 2'-0-methyl
misc_feature
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Sequence 606, Application US/10923330
| Sequence 606, Application US/10923330
| Publication No. US20050153916A1
| APPLICANT: Sirnar Therapeutics, Inc.
| APPLICANT: Sirnar Therapeutics, Inc.
| APPLICANT: McSWAPENTON: Leonid
| TITLE OF INVENTION: Expension Using Short Interfering RNA (sinA)
| FILE REPERENCE: 400/209 (MRH02-708-C)
| CURRENT APPLICATION NUMBER: US/10/223,330
| CURRENT APPLICATION NUMBER: US/204-08-20
| PRIOR PILING DATE: 2004-08-20
| PRIOR PLILING DATE: 2004-06-24
| PRIOR PLILING DATE: 2004-06-24
| PRIOR PLILING DATE: 2004-01-10
| PRIOR APPLICATION NUMBER: PCT/US04/16390
| PRIOR PLILING DATE: 2004-01-16
| PRIOR PLILING DATE: 2003-11-24
| PRIOR PLILING DATE: 2003-11-24
| PRIOR PLILING DATE: 2003-01-36
| PRIOR PLILING DATE: 2003-02-20
| PRIOR PLILING DATE: 2003-03-30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LOCATION: (1). (1)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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NAME/KEY: misc_feature
LOCATION: (3)..(6)
OTHER_INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY: misc_feature
LOCATION: (11)...(11)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
FEATURE:
                                                              148 CCACCGTTCATTCTAGAGC 166
                                                                                                                19 CCACCGTTCATTCTAGAGC 1
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NAME/KEY: misc_feature
LOCATION: (7)...(10)
OTHER INFORMATION: 2'-O-methyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE:
NAME/KEY: misc_feature
LOCATION: (2)...(2)
OTHER INFORMATION: 2'-0-methyl
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OTHER INFORMATION: 2'-O-methyl
FEATURE:
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: misc_feature
LOCATION: (1)...(1)
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OTHER INFORMATION: Description of Artificial Sequence: siNA antisense region
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                PRIOR APPLICATION NUMBER: US 10/826,966
PRIOR FILING DATE: 2004-04-16
PRIOR FILING DATE: 2004-04-16
PRIOR PELING DATE: 2004-04-16
PRIOR PELING DATE: 2004-04-16
PRIOR PELING DATE: 2004-01-124
PRIOR PELING DATE: 2003-11-24
PRIOR APPLICATION NUMBER: US 10/693,059
PRIOR PILING DATE: 2003-11-23
PRIOR PELING DATE: 2003-10-23
PRIOR PELING DATE: 2003-05-23
PRIOR PILING DATE: 2003-05-23
PRIOR APPLICATION NUMBER: PCT/US03/05346
PRIOR PILING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR PILING DATE: 2003-02-20
PRIOR PILING DATE: 2003-02-20
PRIOR PILING DATE: 2003-02-20
PRIOR PILING DATE: 2003-02-20
PREMAIN PRIOR PELING DATE: 2003-02-20
PREMAIN PRIOR PILING DATE: 2003-02-20
PRIOR FILING DATE: 2003-02-20
PRIOR F
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LOCATION: (20) (20)
OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
4.2%; Score 19; DB 1; Length 21;
Best Local Similarity 100.0%; Pred. No. 1e+02;
Matches 19; Conservative 0; Mismatches 0; Indels
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NAME/KEY: misc_feature

COTATION: (21)...(21)

OTHER INFORMATION: n stands for thymidine
US-10-923-330-605
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: misc feature
LOCATION: (1)._(1)
OTHER INFORMATION: 2'-0-methyl
FEATURE:
NAME/KEY: misc feature
LOCATION: (2)._(5)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CATION: (10)...(10)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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NAME/KEY: misc_feature
LOCATION: (14)..(14)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (17)...(17)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
NAME/KEY: misc_feature
LOCATION: (6)._(9)
OTHER INFORMATION: 2'-O-methyl
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LOCATION: (11)...(13)
OTHER INFORMATION: 2'-O-methyl
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LOCATION: (15)...(16)
OTHER INFORMATION: 2'-0-methyl
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LOCATION: (18)...(19)
OTHER INFORMATION: 2'-0-methyl
FILING DATE: 2004-05-24
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LOCATION: (15)..(16)
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JTHER INFORMATION: Description of Artificial Sequence: sinA antisense region
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Sequence 608, Application US/10923330
Publication No. US20050153916A1
GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
APPLICANT: Beigelman, Leonid
APPLICANT: Beigelman, Leonid
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE REPERENCE: 400/209 (MBHB02-708-C)
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT FILING DATE: 2004-08-20
PRIOR FILING DATE: 2004-05-24
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OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
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4.2%; Score 19; DB 1; Length 21;
Best Local Similarity 100.0%; Pred. No. 1e+02;
Matches 19; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: misc_feature; LOCATION: (21)...(21); OTHER INFORMATION: n stands for thymidine US-10-923-330-607
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
NAME/KEX: misc feature
LOCATION: (19) . (19)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                         LOCATION: (4)...(/)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                 LOCATION: (1)._(2)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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                                                                                                                                                                                                                                FEATURE:
NAME/KEX: misc_feature
LOCATION: (3). (3)
OTHER INFORMATION: 2'-0-methyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LOCATION: (17)..(18)
OTHER INFORMATION: 2'-O-methyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY: misc_feature
LOCATION: (8)._(11)
OTHER INFORMATION: 2'-0-methyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LOCATION: (13). (15)
OTHER INFORMATION: 2'-O-methyl
ORGANISM: Artificial Sequence
                                                                                                                                 NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY: misc_feature
LOCATION: (4)..(7)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: misc feature
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LOCATION: (16)..(16)
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APPLICANT: Sirna Therapeutics, Inc.

APPLICANT: McSuiggen, James

APPLICANT: McSuiggen, James

APPLICANT: McSuiggen, James

APPLICANT: Beigelman, Leonid

TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene

TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)

FILE REPRENCE: 400/209 (MSHB2-708-C)

CURRENT ILING DATE: 2004-08-20

FRIOR PRILING DATE: 2004-08-24

PRIOR FILING DATE: 2004-05-24

PRIOR FILING DATE: 2004-05-24

PRIOR FILING DATE: 2004-05-24

PRIOR FILING DATE: 2004-05-46

PRIOR FILING DATE: 2004-01-14

PRIOR FILING DATE: 2004-01-14

PRIOR FILING DATE: 2003-11-24

PRIOR FILING DATE: 2003-10-23

PRIOR FILING DATE: 2003-10-34

PRIOR FILING DATE: 2003-05-23

PRIOR PLILING DATE: 2003-05-23

PRIOR PLILING DATE: 2003-05-36

PRIOR PLILING DATE: 20
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LOCATION: (20) ... (20)
OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 4.2%; Score 19; DB 1; Length 21; Best Local Similarity 100.0%; Pred. No. 1e+02; Matches 19; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: misc_feature

// LOCATION: (21)...(21)

OTHER INFORMATION: n stands for thymidine

US-10-923-330-606
NAME/KEY: misc_feature
LOCATION: (15)...(15)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                   NAME/KEY: misc feature
LOCATION: (18)...(18)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 607, Application US/10923330 publication No. US20050153916A1 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19 CACCGTTCATTCTAGAGCA 1
                                                                                                                                 NAME/KEY: misc feature
LOCATION: (16)...(17)
OTHER INFORMATION: 2'-0-methyl
                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
NAME/KEX: misc_feature
LOCATION: (19) ...(19)
OTHER INFORMATION: 2'-O-methyl
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NAME/KEY: misc feature
LOCATION: (20)...(20)
OTHER INFORMATION: n stands for thymidine
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 ::||||||::||:||
1 CUUCCACCGUUCAUUCUAG 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: RNA
ORGANISM: Artificial Sequence
FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ;
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                               PRIOR FILING DATE: 2002-0-7. Services of PRIOR FILING DATE: 2002-0-7. Services of PRIOR FILING DATE: 2004-0-5.24 PRIOR PELING DATE: 2004-0-5.24 PRIOR APPLICATION NUMBER: PCT/US04/16390 PRIOR PILING DATE: 2004-0-16 PRIOR PAPLICATION NUMBER: US 10/757,803 PRIOR PAPLICATION NUMBER: US 10/757,803 PRIOR PELING DATE: 2004-0-11-24 PRIOR PELING DATE: 2003-11-24 PRIOR PELING DATE: 2003-11-23 PRIOR PELING DATE: 2003-11-23 PRIOR PELING DATE: 2003-11-23 PRIOR PELING DATE: 2003-12-3 PRIOR PELING DATE: 2003-12-3 PRIOR PELING DATE: 2003-12-3 PRIOR PELING DATE: 2003-12-3 PRIOR PELING DATE: 2003-02-20 PRIOR PELING DATE: 2005-02-20 PRIOR PELING D
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NAME/KEY: misc_feature
LOCATION: (20)...(20)
OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
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100.0%; Pred. No. 1e+02;
Live 0; Mismatches 0; Indels
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NAME/KEY: misc_feature
LOCATION: (21)...(21)
OTHER INFORMATION: n stands for thymidine
APPLICATION NUMBER: US 60/396,600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(1)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (4). (4)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (14)...(19)
OTHER_INFORMATION: 2'-deoxy-2'-fluoro
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NAME/KEY: misc_feature
LOCATION: (2)...(3)
OTHER INFORMATION: 2'-O-methyl
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LOCATION: (12)...(13)
OTHER INFORMATION: 2'-O-methyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY: misc_feature
LOCATION: (5)..(8)
OTHER INFORMATION: 2'-0-methyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: RNA
ORGANISM: Artificial Sequence
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Best Local Similarity 100.(
Matches 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FEATURE:
NAME/KEY: misc_feature
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19 GAAGAGTTGGGCTCTGTCA 1

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RESULT 193-

UR-10-923-330-610

Sequence 610, Application US/10923330

Publication No. UG20050153916A1

GENERAL INFORMATION:
APPLICANT: SITUR Therapeutics, Inc.
APPLICANT: Hossiggen, James
APPLICANT: Loonid

TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)

FILE REBERNER: 400/209 (WHH902-708-C)
CURRENT APPLICATION NUMBER: US/10/923,330

CURRENT APPLICATION NUMBER: US/10/923,330

FRIOR PILING DATE: 2004-08-20

PRIOR APPLICATION NUMBER: US/10/926,600

PRIOR APPLICATION NUMBER: US/10/96,600

PRIOR PELLON NUMBER: US/10/96,960

PRIOR PILING DATE: 2004-01-14

PRIOR PILING DATE: 2004-01-14

PRIOR PILING DATE: 2004-01-14

PRIOR PILING DATE: 2003-01-14

PRIOR PILING DATE: 2003-01-14

PRIOR APPLICATION NUMBER: US/10/20,448

PRIOR PILING DATE: 2003-01-24

PRIOR APPLICATION NUMBER: US/10/20,448

PRIOR APPLICATION NUMBER: US/10/20,468

PRIOR APPLICATION NUMBER: US/10/20,468

PRIOR APPLICATION NUMBER: US/10/20,468

PRIOR APPLICATION NUMBER: US/10/20,468

PRIOR PELLING DATE: 2003-02-20

PRIOR PELLOR APPLICATION NUMBER: US/10/20,468

PRIOR PILING DATE: 2003-02-20

REMAINING DATE: 2003-02-
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NAME/KEY: misc_feature
LOCATION: (1)._(1)
OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      . LOCATION: (21)...(21); OTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity US-10-923-330-610
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4.2%; Score 19; DB 1; Length 21;
Best Local Similarity 63.2%; Pred. No. 1e+02;
Matches 12; Conservative 7; Mismatches 0; Indels
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US-10-923-330-611
Sequence 611, Application US/10923330
Publication No. US20050153916A1
GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
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APPLICATION NUMBER: US 60/396,600
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Publication No US20050153916A1
GENERAL INCORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, James
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Expression Using Short Interfering RNA ($1NA)
FILE REFERENCE: 400/209 (MBHB02-708-C)
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT FILING DATE: 2004-08-20
PRIOR FILING DATE: 2004-05-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: Description of Artificial Sequence: sink sense region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FEATURE:
NAMEKEY: misc_feature
LOCATION: (21)...(21)
COTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity
US-10-923-330-611
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LOCATION: (1).7(1)
OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    7; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY: misc feature
LOCATION: (20) ...(20)
OTHER INFORMATION: n stands for thymidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   146 TTCCACCGTTCATTCTAGA 164
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: RNA
ORGANISM: Artificial Sequence
Beigelman, Leonid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity 63.2
Matches 12, Conservative
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NAME/KEY: misc_feature
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OTHER INFORMATION: Description of Artificial Sequence: siNA sense region
PRIOR FILING DATE: 2002-07-17
PRIOR PLICATION NUMBER: PCT/US04/16390
PRIOR PLICATION NUMBER: PCT/US04/16390
PRIOR PLICATION NUMBER: PCT/US04/16390
PRIOR PILING DATE: 2004-06-16
PRIOR PILING DATE: 2004-01-16
PRIOR PLING DATE: 2004-01-14
PRIOR PLING DATE: 2004-01-14
PRIOR PLING DATE: 2003-11-23
PRIOR PLING DATE: 2003-11-23
PRIOR PLING DATE: 2003-11-23
PRIOR PLING DATE: 2003-05-23
PRIOR PLING DATE: 2003-05-23
PRIOR PLING DATE: 2003-02-20
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GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, James
TITLE OF INVENTION: RAN Interference Mediated Inhibition of Telo
TITLE OF INVENTION: RAN Interference Mediated Inhibition of Telo
TITLE OF INVENTION: RAN Interference Mediated Inhibition of Telo
TITLE OF INVENTION: RAN INTERFERENCE: 400/209 (WBHBO2-708-C)
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT APPLICATION NUMBER: PCT/US03/04088
PRIOR FILING DATE: 2004-05-24
PRIOR FILING DATE: 2004-05-24
PRIOR PLICATION NUMBER: PCT/US04/16390
PRIOR FILING DATE: 2004-05-24
PRIOR PLICATION NUMBER: US/05-066
PRIOR FILING DATE: 2004-04-16
PRIOR FILING DATE: 2004-04-16
PRIOR FILING DATE: 2004-04-14
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OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
PEATURE:

j LOCATION: (21)..(21)
j OTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity
US-10-923-330-612

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LOCATION: (20)...(20)
OTHER INFORMATION: n stands for thymidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 613, Application US/10923330 Publication No. US20050153916A1
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 68.41
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
NAME/KEY: misc_feature
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NAME/KEY: misc_feature
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LOCATION:
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US-10-923-330-614

i Sequence 614, Application US/10923330

i Publication No. UG20060153916A1

i GENERAL INFORMATION:
APPLICANT: Sirna Therapeutice, Inc.
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
FILE REFERENCE: 400/209 (MBHB02-708-C)
CURRENT FILING DATE: 2004-08-20

CURRENT FILING DATE: 2004-08-20

FRIOR PILING DATE: 2004-08-24

FRIOR PILING DATE: 2004-07-17

FRIOR APPLICATION NUMBER: US 60/396,600

FRIOR APPLICATION NUMBER: US 60/396,600

FRIOR APPLICATION NUMBER: US 10/4926,966

FRIOR PILING DATE: 2004-01-14

FRIOR FILING DATE: 2004-01-14

FRIOR FILING DATE: 2004-01-14

FRIOR FILING DATE: 2004-01-14

FRIOR FILING DATE: 2003-01-24

FRIOR APPLICATION NUMBER: US 10/593,059

FRIOR APPLICATION NUMBER: US 10/693,059

FRIOR FILING DATE: 2003-01-24

FRIOR FILING DATE: 2003-02-20

FRIOR FILING DATE: 2003-02-20
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PRIOR APPLICATION NUMBER: US 10/720,448

PRIOR FILING DATE: 2003-11-24

PRIOR PELICATION NUMBER: US 10/693,059

PRIOR PELICATION NUMBER: US 10/444,853

PRIOR FILING DATE: 2003-05-23

PRIOR PLING DATE: 2003-05-23

PRIOR PLING DATE: 2003-05-23

PRIOR PLING DATE: 2003-02-20

PRIOR PLING DATE: 2003-02-30

PRIOR PLING DATE: 2003-03-30

PRI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LOCATION: (1).7(1)
OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
4.2%; Score 19; DB 1; Length 21;
Best Local Similarity 73.7%; Pred. No. 1e+02;
Matches 14; Conservative 5; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ION: (20)..(20)
INFORMATION: n stands for thymidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        148 CCACCGTTCATTCTAGAGC 166
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ORGANISM: Artificial Sequence
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NAME/KEY: misc_feature
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LOCATION: (1)...(1)
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OTHER INFO
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PRIOR APPLICATION NUMBER: PCT/USDJ/05028

PRIOR PILOR APPLICATION NUMBER: PCT/USDJ/05028

PRIOR PILOR PLICATION NUMBER: PCT/USDJ/05028

PROBABINED GEOD DAY NOT 12.20.0

PORTER: PROPER NAT 1.20.0

PORTER: PROPER NAT 1.20.0

PROPER NUMBER: PCT/USDJ/05028

PROPER NAT 1.20.0

PROPER NUMBER: PCT/USDJ/05028

PROPER NUMBER: PCT/USDJ/0608

PROPER PROPER PLIAND PARTS: PCT/USDJ/0608

PROPER PLIANT: PLIAND PARTS: 2004-012-22

PROPER PROPER PLIAND PARTS: 2004-012-22

PROPER PLIANT: PURPER PROPER PROPER PLIAND PARTS: PCT/USDJ/0608

PROPER PLIANT: PURPER PROPER PROPE PROPER PROPER
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Best Local Similarity
Matches 19; Conserv
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                                                                      OTHER INFORMATION: Description of Artificial Sequence: sinA sense region
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                                                                                                                                     NAME/KEY: misc_feature
LOCATION: (1)._(1)
OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
FEATURE:
LOCATION: (20)._(20)
OTHER INFORMATION: n stands for thymidine
                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: misc feature
LOCATION: (21) ... (21)
OTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LOCATION: (1). \( (1) \)
OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 19; DB 1; Length 21;
Pred. No. 1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Indels
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 73.7%;
Matches 14; Conservative
ORGANISM: Artificial Sequence
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LOCATION: (1)..(1)
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PUBLICATION NO. US20050153916A1
GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, James
APPLICANT: Baigelman, Leneid
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerage Gene
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerage Gene
TITLE OF INVENTION: Expression Using Short Interfering RNA (8iNA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (8iNA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (8iNA)
CURRENT FILING DATE: 2004-09-20
CURRENT FILING DATE: 2004-05-24
PRIOR APPLICATION WUMBER: PCT/US03/04088
PRIOR PLING DATE: 2004-05-24
PRIOR APPLICATION WUMBER: US 10/826,966
PRIOR PLING DATE: 2004-01-14
PRIOR PLING DATE: 2004-01-14
PRIOR PLING DATE: 2004-01-1-24
PRIOR PLING DATE: 2003-11-24
PRIOR PLING DATE: 2003-11-24
PRIOR PLING DATE: 2003-11-24
PRIOR PLING DATE: 2003-01-34
PRIOR PLING DATE: 2003-02-20
PRIOR APPLICATION WUMBER: US 10/693,059
PRIOR PLING DATE: 2003-02-20
PRIOR APPLICATION WUMBER: US 10/503/05028
PRIOR PLING DATE: 2003-02-20
PRIOR APPLICATION WUMBER: US 10/503/05028
PRIOR PLING DATE: 2003-02-20
PRIOR PLING DATE: 2003-02-20
PRIOR PLING DATE: 2003-02-20
PRIOR PRIOR PRILING DATE: 2003-02-20
PRIOR PRIOR PLING DATE: 2003-02-20
PRIOR PRING PAPLICATION WUMBER: US 10/693,059
PRIOR PLING DATE: 2003-02-20
PRIOR PRING PAPLICATION WUMBER: US 10/693,059
PRIOR PLING DATE: 2003-02-20
PRIOR PRING PAPLICATION WUMBER: US 10/693,059
PRIOR PLING DATE: 2003-03-2-30
PRIOR PLING DATE: 2003-03-3-3
PRIOR PLING PAPLICATION WUMBER: PCT/US03/05028
PRIOR PLING PAPLICATION WUMBER: PCT/US03/05028
PRIOR PLING PAPLICATION PUMBER: PCT/US03/05028
PRIOR PLING PAPLICATION APPLICATION 3-3
PRIOR PLING PAPLICATION APPLICATION 3-3
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OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
NAME/KEY: misc_feature

CCATION: (20)...(20)

COTHER INFORMATION: n stands for thymidine

FEATURE:

NAME/KEY: misc_feature

LOCATION: (21)...(21)

COTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity
US-10-923-330-616
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                                                                                                                                                                                                                                                                                                                                                            Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                            Query Match
4.2%; Score 19; DB 1;
Best Local Similarity 73.7%; Pred. No. 1e+02;
Matches 14; Conservative 5; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PEATURE:

VAME/KEY: misc_feature

LOCATION: (21)...(21)

OTHER INFORMATION: n stands for thymidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 618, Application US/10923330 Publication No. US20050153916A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              300 GAAGAGTTGGGCTCTGTCA 318
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US-10-923-330-621/c
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APPLICANT: Beigelman, James James
APPLICANT: Beigelman, James
APPLICANT: Beigelman, James
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
CURRENT APPLICATION NUMBER: PCT/US03/0408
PRIOR PILING DATE: 2004-05-24
PRIOR PELING DATE: 2004-05-24
PRIOR PELING DATE: 2004-05-24
PRIOR PILING DATE: 2004-05-24
PRIOR PILING DATE: 2004-01-14
PRIOR PLING DATE: 2004-01-14
PRIOR PLING DATE: 2003-11-23
PRIOR PLING DATE: 2003-11-24
PRIOR PLING DATE: 2003-01-24
PRIOR PLING DATE: 2003-01-24
PRIOR PLING DATE: 2003-02-20
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SOFTWARE: Patentin version 3.3
SEQ ID NO 619
LENGTH: 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              .;
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4.2%; Score 19; DB 1; Length 21;
Best Local Similarity 100.0%; Pred. No. 1e+02;
Matches 19; Conservative 0; Mismatches 0; Indels
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NAMB/KEY: misc feature
LOCATION: (21)
COTHER INFORMATION: n stands for thymidine
US-10-923-330-619
                                                                                                                                                                                                              Sequence 619, Application US/10923330
Publication No. US20050153916A1
GENERAL INFORMATION:
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145 CTTCCACCGTTCATTCTAG 163
                                      19 CTTCCACCGTTCATTCTAG 1
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ORGANISM: Artificial Sequence
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LOCATION: (20)..(20)
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US-10-923-330-620/c
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APPLICANT: Resulgen, James
TITLE OF INVENTION: Resulgen, James
TITLE OF INVENTION: Repression Using Short Interfering RNA (sita)
TITLE OF INVENTION: RAY INTERference Mediated Inhibition of Tollamerase Gene
TITLE OF INVENTION: RAY INTERference Mediated Inhibition of Tollamerase Gene
TITLE OF INVENTION: RAY INTERFERENCE: 40/0.20 (WEBBIO.708-0.0)
FRICA APPLICATION WINDER: US/0.20-0.3.30
PRICA APPLICATION WINDER: US/0.20-3.30
PRICA APPL
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JS-10-923-330-623/c
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US-10-923-330-622/C

i Sequence 622, Application US/10923330

i Publication No. US20060153016A1

i GENERAL INFORMATION:

i APPLICANT: Sirna Therapeutics, Inc.

i APPLICANT: McSwiggan, James

i TILLE OF INVENTION: Expression Using Short Interfering RNA (siNA)

FILE REFREENCE: 400/209 (MBHBR2-708-C)

CURRENT APPLICATION NUMBER: US/10/923,330

CURRENT FILING DATE: 2004-06-24

PRIOR FILING DATE: 2004-05-24

PRIOR APPLICATION NUMBER: US 60/396,600

PRIOR PILING DATE: 2004-01-14

PRIOR APPLICATION NUMBER: US 10/720,448

PRIOR APPLICATION NUMBER: US 10/444,853

PRIOR APPLICATION NUMBER: US 10/444,853

PRIOR APPLICATION NUMBER: US 10/444,853

PRIOR APPLICATION NUMBER: US 10/309650

PRIOR APPLICATION NUMBER: US 10/309650

PRIOR APPLICATION NUMBER: US 10/309650

PRIOR APPLICATION NUMBER: US 10/444,853

PRIOR APPLICATION NUMBER: US 10/444,853
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PRIOR FILING DATE: 2004-05-24
PRIOR PLICATION NUMBER: US 10/826,966
PRIOR FILING DATE: 2004-01-16
PRIOR FILING DATE: 2004-01-14
PRIOR PLICATION NUMBER: US 10/757,803
PRIOR PLICATION NUMBER: US 10/720,448
PRIOR PLICATION NUMBER: US 10/720,448
PRIOR FILING DATE: 2003-11-23
PRIOR PLICATION NUMBER: US 10/693,059
PRIOR PLICATION NUMBER: US 10/444,853
PRIOR PLILING DATE: 2003-05-23
PRIOR PLILING DATE: 2003-05-20
PRIOR PLILING DATE: 2003-02-20
PRIOR PLILING DATE: 2003-02-20
PRIOR PLILING DATE: 2003-02-20
PRIOR FILING DATE: 2003-02-20
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COATION: (20)...(20)

OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
FRATURE:
NAME/KEY: misc_feature
LOCATION: (21)...(21)

COTHER INFORMATION: 010...(21)

US-10-923-330-621
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100.0%; Pred. No. 1e+02;
ive 0; Mismatches 0; Indels
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ORGANISM: Artificial Sequence
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Best Local Similarity
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FEATURE: OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region
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APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
APPLICANT: Beigelman, Leoner
APPLICANT: Beigelman, Leoner
APPLICANT: Beigelman, Leoner
APPLICANT: Beigelman, Leoner
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION NUMBER: US/10/923,330
CURRENT FILING DATE: 2004-08-20
PRIOR APPLICATION NUMBER: PCT/US03/04088
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gapa
PRIOR FILING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR FILING DATE: 2003-02-20
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 768
SOFTWARE: PatentIn version 3.3
SEQ ID NO 622
LENGTH: 21
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NAME/KEY: misc_feature
LOCATION: (20)...(20)
OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
FEATURE:
NAME/KEY: misc_feature
NAME/KEY: misc_feature
LOCATION: (21)...(21)
CTHER INFORMATION: n stands for thymidine
US-10-923-330-622
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 21;
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1e+02;
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PRIOR PLING DATE: 2004-05-24
PRIOR PLING DATE: 2006-07-17
PRIOR PELING DATE: 2004-05-27
PRIOR APPLICATION NUMBER: US 10/720,416390
PRIOR PLING DATE: 2004-05-16
PRIOR PLING DATE: 2004-01-14
PRIOR PLING DATE: 2004-01-14
PRIOR PLING DATE: 2003-11-24
PRIOR APPLICATION NUMBER: US 10/720,448
PRIOR PLING DATE: 2003-11-24
PRIOR PLING DATE: 2003-11-23
PRIOR PLING DATE: 2003-11-23
PRIOR PLING DATE: 2003-11-23
PRIOR PLING DATE: 2003-11-23
PRIOR PLING DATE: 2003-10-33
PRIOR PLING DATE: 2003-10-33
PRIOR PLING DATE: 2003-05-20
PRIOR PLING DATE: 2003-02-20
PRIOR PLING DATE: 2003-02-20
PRIOR PLING DATE: 2003-02-20
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                                                                                                                                                                                                                                                                          TYPE: RNA
ORGANISM: Artificial Sequence
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US-10-923-330-626/c
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                                     Matches
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WESULY 2013

WESULY 2020

WESULY 2020

WESULY 2020

WESULY 2020

SEQUENCE 624, Application US/10923330

Publication No. US20050153916A1

GENERAL INFORMATION:

MESWIGHOLD AND THE PROPERLY SITUED

APPLICANT: Being Therepeutics, Inc.

APPLICANT: Being Properties, Inc.

APPLICANT: Being Properties and Thereference Mediated Inhibition of Telomerase Gene TITLE OF INVENTION: RAY Interference Mediated Inhibition of Telomerase Gene TITLE OF INVENTION: RAPES 100/209

TITLE OF INVENTION: RAPES: 2004-02-0

CURRENT APPLICATION NUMBER: DET/1092/3,330

PRIOR APPLICATION NUMBER: US 60/396,600

PRIOR FILING DATE: 2004-05-24

PRIOR PELING DATE: 2002-07-17

PRIOR PELING DATE: 2004-04-16

PRIOR APPLICATION NUMBER: US 10/757,803

PRIOR APPLICATION NUMBER: US 10/757,803

PRIOR APPLICATION NUMBER: US 10/757,803

PRIOR PELING DATE: 2003-11-23

PRIOR PILING DATE: 2003-11-23

PRIOR PELING DATE: 2003-11-23

PRIOR PILING DATE: 2003-11-23

PRIOR PELING DATE: 2003-11-23

PRIOR PILING DATE: 2003-11-23

PRIOR PELING DATE: 2003-12-20

PRIOR PELING DATE: 2003
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LOCATION: (21)..(21)

// OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage

US-10-923-330-623
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OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
                                                                                                                                                                                                                                                                                                                                                                    DB 1; Length 21;
1e+02;
hes 0; Indels
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Best Local Similarity 100.0%; Pred. No. 1e+02;
                                                                                                                                                                                                                                                                                                                                                            Query Match 4.2%; Score 19; DB Best Local Similarity 100.0%; Pred. No. leamatches 19; Conservative 0; Mismatches
FEATURE:
NAME/KEY: misc_feature
NOCATION: (20)...(20)
OTHER INFORMATION: n stands for thymidine
FEATURE:
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; OTHER INFORMATION: n stands for thymidine
US-10-923-330-624
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       150 ACCGTTCATTCTAGAGCAA 168
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LOCATION: (20)..(20)
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Sequence 626, Application US/10923330
| Sequence 626, Application US/10923330
| Publication No. US20050153916A1
| GENERAL INFORMATION:
| APPLICANT: Sitna Therapeutics, Inc.
| APPLICANT: MoSWAGEN, James
| APPLICANT: MOSWAGEN, James
| APPLICANT: MOSWAGEN, James
| APPLICANT: MOSWAGEN, James
| APPLICANT: Beigelman, Leonid
| TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
| FILE REPERENCE: 400/209 (MRH02-708-C)
| CURRENT PAPLICATION NUMBER: US/10/923,330
| CURRENT PELLING DATE: 2004-06-20
| PRIOR PILING DATE: 2004-06-20
| PRIOR PELLOR DATE: 2004-06-20
| PRIOR PELLOR DATE: 2004-06-10
| PRIOR PELLOR DATE: 2004-06-10
| PRIOR PELLOR DATE: 2004-01-10
| PRIOR PELLOR DATE: 2004-01-11
| PRIOR PELLOR DATE: 2004-01-11
| PRIOR PELLOR DATE: 2004-01-12
| PRIOR PELLOR DATE: 2004-01-12
| PRIOR PELLOR DATE: 2004-01-12
| PRIOR PELLOR DATE: 2003-10-23
| PRIOR PELLOR DATE: 2003-01-23
| PRIOR PELLOR DATE: 2003-02-20
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               Gaps
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       0; Mismatches
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NAME/KEY: misc_feature
LOCATION: (1)..(2)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (7). (7)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (3)...(6)
OTHER INFORMATION: 2'-O-methyl
FBAUURE:
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LOCATION: (8). (10)
OTHER INFORMATION: 2'-O-methyl
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LOCATION: (12)...(13)
OTHER INFORMATION: 2'-0-methyl
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ORGANISM: Artificial Sequence
FEATURE:
Conservative
                                                                      300 GAAGAGTIGGGCTC
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US-10-923-330-628/c
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US-10-923-330-627/A

Sequence 637, Application US/10923330

PUBLIcation No. US20050153916A1

GENERAL INFORMATION:

APPLICANT: SITEMA Therapeutics, Inc.

APPLICANT: MCSWAJGEN, James

APPLICANTON: Expression Using Short Interfering RNA (sinA)

FILE OF INVENTION: Expression Using Short Interfering RNA (sinA)

FILE OF INVENTION: Expression Using Short Interfering RNA (sinA)

FILE OF INVENTION WUBBER: US/10/923,330

CURRENT FILING DATE: 2004-08-24

FRIOR APPLICATION WUBBER: US 0/396,600

FRIOR PLICATION WUBBER: US 10/826,966

FRIOR PLICATION WUBBER: US 10/826,966

FRIOR APPLICATION WUBBER: US 10/757,803

FRIOR APPLICATION WUBBER: US 10/757,448

FRIOR PLICATION WUBBER: US 10/444,853

FRIOR PLILING DATE: 2003-11-23

FRIOR PLILING DATE: 2003-11-23

FRIOR PLILING DATE: 2003-11-23

FRIOR PLILING DATE: 2003-11-23

FRIOR PLILING DATE: 2003-10-20

FRIOR APPLICATION WUBBER: PCT/US03/05028

FRIOR PLILING DATE: 2003-05-23

FRIOR PLILING DATE: 2003-05-23

FRIOR APPLICATION WUBBER: PCT/US03/05028

FRIOR PLILING DATE: 2003-05-23

FRIOR APPLICATION WUBBER: PCT/US03/05028

FRIOR PLILING DATE: 2003-05-23

FRIOR PLILING DATE: 2003-05-23

FRIOR APPLICATION WUBBER: PCT/US03/05028

FRIOR PLING DATE: 2003-05-23

FRIOR PLING APPLICATION WUBBER: PCT/US03/05028

FRIOR PLING APPLICATION WUBBER: PCT/US03/05028

FRIOR PLING APPLICATION WUBBER: 
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| LOCATION: (21)...(21)

| OTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity

US-10-923-330-626
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Pred. No. 1e+02;
0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                   FEATURE:
NAME/KEY: misc feature
LOCATION: (20) ...(20)
OTHER INFORMATION: n stands for thymidine
NAME/KEY: misc_feature
LOCATION: (14) ... (14)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: misc_feature
LOCATION: (1)._(3)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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                                                                                                  FEATURE:
NAME/KEY: misc_feature
LOCATION: (15)..(19)
OTHER INFORMATION: 2'-0-methyl
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ORGANISM: Artificial Sequence
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US-10-924-310-628(Application US/10923330)

| US-10-924-310-628(Application US/10923330)
| Publication No. US20050153916A1 |
| GENERAL INFORMATION: |
| APPLICANT: Sirna Therapeutics, Inc. |
| APPLICANT: Beigelman, James |
| APPLICANT: Beigelman, Leonid |
| TITLE OF INVENTION: Expression Using Short Interfering RNA (BiNA) |
| TITLE OF INVENTION: Expression Using Short Interfering RNA (BiNA) |
| TITLE OF INVENTION: Expression Using Short Interfering RNA (BiNA) |
| TITLE OF INVENTION: Expression Using Short Interfering RNA (BiNA) |
| FILE REPRENENCE: 4006/209 (MBHB02-708-C) |
| FRIOR FILING DATE: 2004-08-20 (Bina) |
| FRIOR FILING DATE: 2004-05-24 (Bina) 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gapa
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OTHER INFORMATION: n stands for thymidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: misc feature
LOCATION: (12)...(12)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                             LOCATION: (8). (8)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (4).7(7)
OTHER INFORMATION: 2'-0-methyl
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OTHER INFORMATION: 2'-0-methyl
                                                                                                                                                                                                                                                                                   NAME/KEY: misc_feature
LOCATION: (9)._(11)
OTHER INFORMATION: 2'-0-methyl
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OTHER INFORMATION: 2'-0-methyl
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Best Local Similarity 100.0
Matches 19; Conservative
                                                             FEATURE:
NAME/KEY: misc_feature
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FEATURE:
NAME/KEY: misc_feature
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PRIOR FILING DATE: 2003-11-23
PRIOR APPLICATION NUMBER: US 10/444,853
PRIOR FILING DATE: 2003-05-23
PRIOR PELING DATE: 2003-05-22
PRIOR APPLICATION NUMBER: PCT/US03/05346
PRIOR PILING DATE: 2003-02-20
PRIOR PILING DATE: 2003-02-20
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ 1D NOS: 768
SOFTWARE: PatentIn version 3.3
SEQ ID NO 628
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NAME/KEY: misc feature

LOCATION: (21)

CTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity
US-10-923-330-628
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100.0%; Pred. No. 1e+02;
ive 0; Mismatches 0; Indels
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OTHER INFORMATION: n stands for thymidine
                                                                                                                                                                                                                                                                                                                                                                                                                  LOCATION: (1). (4)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: misc feature
LOCATION: (9)._(9)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LOCATION: (13)...(13)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (16)...(16)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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US-10-923-330-629/c
Sequence 629, Application US/10923330
Publication No. US20050153916A1
GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   147 TCCACGTTCATTCTAGAG 165
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LOCATION: (5). (8)
OTHER INFORMATION: 2'-0-methyl
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LOCATION: (10)...(12)
OTHER INFORMATION: 2'-0-methyl
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LOCATION: (14)...(15)
OTHER INFORMATION: 2'-0-methyl
                                                                                                                                                                                                                                                                       TYPE: RNA
ORGANISM: Artificial Sequence
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OTHER INFORMATION: 2'-0-methyl
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Best Local Similarity 100.
Matches 19; Conservative
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LOCATION: (1)..(4)
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NAME/KEY: misc feature
LOCATION: (13)...(13)
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LOCATION: (17)..(19)
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LOCATION: (20)...(20)
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APPLICANT: Websigen, James
APPLICANT: Websigen, James
TITLE OF INDEPTIBATION Expression Using Abdiated Inhibition of Telemerase Gene
TITLE OF INDEPTIBATION EXPRESSION USING ABDIATED THE OF INDEPTIBATION EXPRESSION USING ABDIATED THE OF INDEPTIBATION EXPRESSION USING ABDIATED THE OF INDEPTIBATION OF THE OFFICE THE OFFI
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US-10-923-310-530.

US-10-923-310-530.

US-10-923-310-930.

US-10-923-310-930.

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                                                                                                           Query Match

4.2%; Score 19; DB 1; Length 21;
Best Local Similarity 100.0%; Pred. No. 1e+02;
Matches 19; Conservative 0; Mismatches 0; Indels
NAME/KEY: misc feature
LOCATION: (20) ... (20)
OTHER INFORMATION: n stands for thymidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FEATURE:
NAME/KEY: misc_feature
LOCATION: (3)._(6)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
FEATURE:
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NAME/KEX: misc_feature
LOCATION: (1): (1)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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NAME/KEXY: misc_feature
LOCATLON: (2)..(2)
OTHER INFORMATION: 2'-O-methyl
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ORGANISM: Artificial Sequence
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US-10-923-330-630/c
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GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Meswiggen, James
APPLICANT: Meswiggen, James
APPLICANT: Meswiggen, James
TITLE OF INVENTION: RAD Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RAD INTERFORMER: AD0/209 (MBHB02-708-C)
CURRENT APPLICATION NUMBER: PG/10/923,330
CURRENT FILING DATE: 2004-08-20
FRIOR FILING DATE: 2004-05-24
FRIOR FILING DATE: 2004-05-24
FRIOR APPLICATION NUMBER: US 10/826,966
FRIOR FILING DATE: 2004-04-16
FRIOR FILING DATE: 2004-04-16
FRIOR PELING DATE: 2004-04-16
FRIOR PELING DATE: 2004-01-14
FRIOR PELING DATE: 2004-01-14
FRIOR PELING DATE: 2004-01-14
FRIOR FILING DATE: 2004-01-14
FRIOR FILING DATE: 2004-01-14
FRIOR FILING DATE: 2003-11-24
FRIOR FILING DATE: 2003-11-24
FRIOR FILING DATE: 2003-11-23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME/KEY: misc feature
LOCATION: (20)...(20)
OTHER INFORMATION: n stands for thymidine
                                                                                                                                                                                                                                                                                                                                                   LOCATION: (15)...(15)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: misc_feature
LOCATION: (18)..(18)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
FEATURE:
                                                                                                                                LOCATION: (11)...(11)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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; Publication No. US20050153916A1
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                                                                                                                                                                                FEATURE:
NAME/KEX: misc_feature
LOCATION: (12)..(14)
OTHER INFORMATION: 2'-0-methyl
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LOCATION: (16)..(17)
OTHER INFORMATION: 2'-O-methyl
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LOCATION: (19)..(19)
OTHER INFORMATION: 2'-0-methyl
                                             OTHER INFORMATION: 2'-O-methyl
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NAME/KEY: misc_feature
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NAME/KEY: misc_feature
NAME/KEY: misc_feature
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Publication No. US20050153916A1
                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: Description of Artificial Sequence: siNA antisense region
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PRIOR APPLICATION NUMBER: US 10/444,853
PRIOR FILING DATE: 2003-05-23
PRIOR APPLICATION NUMBER: PCT/US03/05346
PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR PILING DATE: 2003-02-20
PRIOR FILING PAID: 2003-02-20
Remaining Prior Application data removed - See File Wrapper or PALM.
SOFTWARE: Patentin version 3.3
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NAME/KEY: misc_feature
LOCATION: (21)...(21)
OTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity
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4.2%; Score 19; DB 1; Length 21;
Best Local Similarity 100.0%; Pred. No. 1e+02;
Matches 19; Conservative 0; Mismatches 0; Indels
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NAME/KEY: misc_feature
LOCATION: (20)...(20)
OTHER INFORMATION: n stands for thymidine
                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY: misc_feature
LOCATION: (1)._(2)
OTHER_INFORMATION: 2'-deoxy-2'-fluoro
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INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (12)...(12)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (16)...(16)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (19)...(19)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (3)...(3)
OTHER INFORMATION: 2'-O-methyl
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NAME/KRY: misc feature
LOCATION: (8). (11)

OTHER INFORMATION: 2'-0-methyl
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NAME/KEY: misc_feature
LOCATION: (13)..(15)
OTHER INFORMATION: 2'-0-methyl
                                                                                                                                                                                                                                                       TYPE: RNA
ORGANISM: Artificial Sequence
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LOCATION: (17)..(18)
OTHER INFORMATION: 2'-O-methyl
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                                                                                                                                                                                                            SEQ ID NO 631
LENGTH: 21
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RESULT 213 US-10-923-330-632/c ; Sequence 632, Application US/10923330

150 ACCGTTCATTCTAGAGCAA 168

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19 ACCGTTCATTCTAGAGCAA 1

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APPLICANT NO. 12500513315A1

APPLICANT NO. 1250051305A1

TITLE OF INVENTION: RNA INTERFERENCE ANGLE AND THE APPLICANT NO. 125005A1

TITLE OF INVENTION: RNA INTERFERENCE ANGLE ANG
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                                                                                                                                  OTHER INFORMATION: Description of Artificial Sequence: siNA antisense region
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                                                                 Gaps
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; LOCATION: (21)...(21)
; OTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity US-10-923-330-632
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; OTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity
US-10-923-330-634
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                                          4.2%; Score 19; DB 1; Length 21;
100.0%; Pred. No. 1e+02;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: misc feature
LOCATION: (20) ... (20)
OTHER INFORMATION: n stands for thymidine
                                                                                   300 GAAGAGTTGGGCTCTGTCA 318
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         145 CTTCCACCGTTCATTCTAG 163
                                                                                                        19 GAAGAGTTGGGCTCTGTCA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: RNA
ORGANISM: Artificial Sequence
                                        Query Match 4.2%;
Best Local Similarity 100.0%
Matches 19; Conservative
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19 CTTCCACCGTTCATTCTAG 1

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RESULT. 15.0, 438/C.

RESULT. 15.0, 438/C.
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OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region
              PRIOR APPLICATION NUMBER: US 10/720,448

PRIOR FILING DATE: 2003-11-24

PRIOR PLILING DATE: 2003-11-24

PRIOR PLILING DATE: 2003-11-24

PRIOR PLILING DATE: 2003-11-23

PRIOR PLILING DATE: 2003-05-23

PRIOR PLILING DATE: 2003-05-23

PRIOR PLILING DATE: 2003-02-20

PRIOR PLILING DATE: 2003-02-20

PRIOR PLILING DATE: 2003-02-20

PRIOR PLILING DATE: 2003-02-20

Remaining Prior Application data removed - See File Wrapper or PALM.

SOFTWARE: Patentin version 3.3

LENGTH: 21

LENGTH: 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:

NAME/KEY: misc_feature

LOCATION: (20)...(20)

CTHER INFORMATION: n stands for thymidine

FEATURE:

NAME/KEY: misc_feature

LOCATION: (21)...(21)

CTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity

US-10-923-330-637
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100.0%; Pred. No. 100.0%; 0; Mismatches
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Best Local Similarity 100.0
Matches 19; Conservative
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| Publication No. US20050153916A1
| GENERAL INFORMATION:
| APPLICANT: Sirna Therapeutics, Inc.
| APPLICANT: Sirna Therapeutics, Inc.
| APPLICANT: Beigalman, Leonid
| TITLE OF INVENTION: Expression Using Short Interfering RNA (sina)
| TITLE OF INVENTION: Expression Using Short Interfering RNA (sina)
| TITLE OF INVENTION: Expression Using Short Interfering RNA (sina)
| TITLE OF INVENTION: Expression Using Short Interfering RNA (sina)
| TITLE OF INVENTION: Expression Using Short Interfering RNA (sina)
| TITLE OF INVENTION: Expression Using Short Interfering RNA (sina)
| TITLE OF INVENTION: Expression Using Short Interfering RNA (sina)
| TITLE OF INVENTION: Expression Using Short Interfering RNA (sina)
| FRICA FILLING DATE: 2004-06-20
| PRIOR FILLING DATE: 2004-06-20
| PRIOR FILLING DATE: 2004-06-24
| PRIOR FILLING DATE: 2004-04-16-65-34
| PRIOR FILLING DATE: 2004-04-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region
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                      CURRENT FILING DATE: 2004-08-20

PRIOR PELING DATE: 2004-08-20

PRIOR PILING DATE: 2004-06-24

PRIOR FILING DATE: 2004-06-24

PRIOR FILING DATE: 2004-06-136,600

PRIOR PELING DATE: 2004-05-17

PRIOR APPLICATION NUMBER: US 10/826,966

PRIOR PILING DATE: 2004-04-16

PRIOR PILING DATE: 2004-04-14

PRIOR PILING DATE: 2004-04-14

PRIOR PILING DATE: 2003-11-4

PRIOR APPLICATION NUMBER: US 10/720,448

PRIOR APPLICATION NUMBER: US 10/693,059

PRIOR APPLICATION NUMBER: US 10/693,059

PRIOR PILING DATE: 2003-11-23

PRIOR PILING DATE: 2003-11-23

PRIOR PILING DATE: 2003-02-20

PRIOR PILING DATE: 2003-02-20
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4.2%; Score 19; DB 1; Length 21;
Best Local Similarity 100.0%; Pred, No. 1e+02;
Matches 19; Conservative 0; Mismatches 0; Indels
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LOCATION: (20) ... (20)
OTHER INFORMATION: n stands for thymidine
CURRENT APPLICATION NUMBER: US/10/923,330
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: RNA
ORGANISM: Artificial Sequence
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LOCATION: (21)..(21)
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LENGTH: 21
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Gaps

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; NAME/KEY: misc_feature
; LOCATION: (21)
; OTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity
US-10-923-330-639
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Best Local Similarity
Matches 19; Conserv
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US-10-923-330-640/c
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Sequence 639, Application US/10923330

Publication No. US20050153916A1

APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: RAN Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RAN Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE REFERENCE: 400/209 (MBHB02-708-C)
CURRENT APPLICATION NUMBER: US/10/923,330

CURRENT FILING DATE: 2004-05-24

PRIOR FILING DATE: 2004-05-24

PRIOR PELING DATE: 2004-05-24

PRIOR PELING DATE: 2004-01-14

PRIOR APPLICATION NUMBER: US 10/826,966

PRIOR APPLICATION NUMBER: US 10/826,966

PRIOR APPLICATION NUMBER: US 10/757,803

PRIOR APPLICATION NUMBER: US 10/757,803

PRIOR APPLICATION NUMBER: US 10/757,803

PRIOR PELING DATE: 2003-11-24

PRIOR PELING DATE: 2003-12-20

PRIOR PELING DATE: 2003-13-24

PRIOR PELING DATE: 2003-13-34

PRIOR PELING DATE: 2003-02-20

PRIOR PELING DATE: 2003-03-23

PRIOR PELING DATE: 2003-0
                                                                                                                                                                                                                        OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region
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                                                                                                                                                                                                                                                  FEATURE:
NAME/KEY: misc_feature
LOCATION: (20)..(20)
OTHER INFORMATION: n stands for thymidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: misc_feature
LOCATION: (20)..(20)
OTHER INFORMATION: n stands for thymidine
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 149 CACCGTTCATTCTAGAGCA 167
   SOFTWARE: PatentIn version 3.3 SEQ ID NO 638 LENGTH: 21
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ORGANISM: Artificial Sequence
                                                                                                                                                 ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY: misc feature
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US-10-923-330-639/c
                                                                                                                   TYPE: RNA
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GENERAL INFORMATION:
JAPRICANT: Sirna Therapeutics, Inc.
APPLICANT: McSWaggen, James
APPLICANT: McSWaggen, James
APPLICANT: McSWaggen, James
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
FILE REPERENCE: 400/209 (WHBH02-708-C)
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT APPLICATION NUMBER: PCT/US04/1639
PRIOR FILING DATE: 2004-06-24
PRIOR FILING DATE: 2004-05-24
PRIOR FILING DATE: 2004-05-24
PRIOR PLILING DATE: 2004-01-14
PRIOR PLILING DATE: 2004-05-23
PRIOR PLILING DATE: 2004-05-23
PRIOR PLILING DATE: 2003-05-23
PRIOR PLILING DATE: 2003-05-24
PRIOR PLILING DATE: 2003-05-24
PRIOR PLILING DATE: 2003-05-24
PRIOR PLILING DATE: 2003-05-25
PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR PLILING DATE: 2003-05-24
PRIOR PLILING DATE: 2003-05-26
PRIOR PLILING DATE: 2003-05-27
PRIOR PRIOR PLILING DATE: 2003-05-27
PRIOR PRIOR PLILING DATE: 2003-05-27
PRIOR PRIOR PLILING DATE: 2003-05-36
PRIOR PLILING DATE: 2003-05-36
PRIOR PLILING DATE: 2003-05-37
PRIOR
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1 LOCATION: (21)...(21)

2 OTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity

US-10-923-330-640
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Length 21;
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4.2%; Score 19; DB 1;
Best Local Similarity 100.0%; Pred. No. 1e+02;
Matches 19; Conservative 0; Mismatches 0
ch 4.2%; Score 19; DB 1; 1 Similarity 100.0%; Pred. No. 1e+02; 19; Conservative 0; Mismatches (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: misc_feature
LCCATION: (20)...(20)
OTHER INFORMATION: n stands for thymidine
PEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 640, Application US/10923330 Publication No. US20050153916Al GENERAL INFORMATION:
                                                                                                                                                                 150 ACCGTTCATTCTAGAGCAA 168
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                                                                                                                                                                                                               19 ACCGTTCATTCTAGAGCAA 1
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                                                                                  APPLICANT: Kilian, Andrzej
APPLICANT: Kilian, Andrzej
APPLICANT: Bowtell, David
TITLE OP INVENTION: VERTEBRATE TELOMERASE GENES AND PROTEINS AND USES
TITLE OP INVENTION: THEREOF
FILE REFERENCE: TWGL 40723
CURRENT APPLICATION NUMBER: US/10/900,231
CURRENT FILING DATE: 2004-07-27
NUMBER OF SEO ID NOS: 155
SOFTWARE: PATENTIN Ver. 2.0
SEO ID NO 126
LENGTH: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: Description of Artificial Sequence: Synthesized OTHER INFORMATION: Amplification Primer Design based on EST Sequence US-10-900-231-126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           .;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 543, Application US/10325810
| Publication No. US20030204069A1
| GENERAL INFORMATION:
| Lingner, Joachim | Nakamura, Toru | Chapman, Karen B. | Morin, Gregg B. | Harley, Calvin B. | Andrews, William H. | TITLE OF INVENTION: Walliam H. | TITLE OF INVENTION: Human Telomerase Catalytic Subunit | NUMBER OF SEQUENCES: 633
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4.0%; Score 18; DB 1; Length 18;
100.0%; Pred. No. 93;
iive 0; Mismatches 0; Indels
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION DATA:

APPLICATION DATA:

APPLICATION NUMBER: US/10/325,810

FILING DATE: 20-Dec-2002

CLASSIFICATION: UNKNOWN-

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/09/402,181

FILING DATE: 29-SEP-1997

APPLICATION NUMBER: US 08/724,643

FILING DATE: 01-OCT-1996

APPLICATION NUMBER: US 08/844,419

FILING DATE: 18-APR-1997

APPLICATION NUMBER: US 08/844,419

FILING DATE: 18-APR-1997
                      Sequence 126, Application US/10900231
Publication No. US20050176022A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 GGGTTGCGGAGGGTGGGC 18
                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 GGGTTGCGGAGGGTGGGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 100.
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 222
US-10-325-810-543/c
-10-900-231-126
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Gaps
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| Sequence 543, A4plication US/10877124
| Publication No. US2004024529A1
| GENERAL INFORMATION:
| Lingner, Joachim | Lingner, Joachim | Makamura, Toru | Chapman, Karen B. | Morin, Greeg B. | Harley, Calvin B. | Andrews, William H. |
| TITLE OF INVENTION: Human Telomerase Catalytic Subunit NUMBER OF SEQUENCES: 727 | CORRESPONDENCE ADDRESS: 727 | CORRESPONDENCE ADDRESS: ADDRESSE: Two Embarcadero Center, Eighth Floor | CITY: San Francisco | STATE: California | COUNTRY: USA | COMPUTER READABLE FORM: | MEDIUM TYPE: TWO FORM: | MEDIUM TYPE: |
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COCATION: 1..18
OTHER INFORMATION: /note= "antisense hTRT molecule"
SEQUENCE DESCRIPTION: SEQ ID NO: 543:
US-10-325-810-543
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3.6%; Score 16.4; DB 1; Length 18; 94.4%; Pred. No. 1.4e+02; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/877,124
FILING DATE: 24-Jun-2004
CLASSIFICATION: <UNknown>
                                                                                                                                                                                                                                                                                 FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/912,951
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: WO 0F/1/US97/17885
FILING DATE: 01-OCT-1997
ATTORNEY/AGENT INFORMATION:
NAME: Ausenhus, Scott L.
REGISTRATION NUMBER: 42,271
REFERENCE/DOCKET NUMBER: 015389-002620US
TELECOMMUNICATION INFORMATION:
                               FILING DATE: 06-MAY-1997
APPLICATION NUMBER: US 08/854,050
FILING DATE: 09-MAY-1997
APPLICATION NUMBER: US 08/911,312
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/912,951
APPLICATION NUMBER: US 08/851,843
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (415) 576-0300 INFORMATION FOR SEQ ID NO: 543: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             149 CACCGTTCATTCTAGAGC 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 18 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18 caccerrearreragage 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 3.6
Best Local Similarity 94.4
Matches 17; Conservative
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US-10-877-124-543/c
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Gaps
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OTHER INFORMATION: /note= "antisense hTRT molecule"
SEQUENCE DESCRIPTION: SEQ ID NO: 543:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 3.6%; Score 16.4; DB 1; Length 18; Best Local Similarity 94.4%; Pred. No. 1.4e+02; Matches 17; Conservative 0; Mismatches 1; Indels
                                                                                                                       SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/877,022
FILING DATE: 24-Jun-2004
CLASSIFICATION: <UNknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/912,951
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/915,503
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: WO PCT/US97/17618
FILING DATE: 01-OCT-1997
APPLICATION NUMBER: WO PCT/US97/17885
FILING DATE: 01-OCT-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Apple, Randolph Ted
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002610US
                                                                                                                                                                                                                                                                                             FILING DATE: 02-Nov-1999
APPLICATION NUMBER: 08/974,549
FILING DATE: 4Unknown>
APPLICATION NUMBER: 08/844,419
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/846,017
FILING DATE: 25-APR-1997
APPLICATION NUMBER: US 08/851,843
FILING DATE: 06-MAY-1997
APPLICATION NUMBER: US 08/854,050
FILING DATE: 09-MAY-1997
APPLICATION NUMBER: US 08/854,050
FILING DATE: 09-MAY-1997
APPLICATION NUMBER: US 08/854,050
                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/432,503
ZIP: 94111-3834
COMPUTER READMBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 543:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 543, Application US/10877146 Publication No. US20050013825A1 GENERAL INFORMATION:
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Lingner, Joachim
Nakamura, Toru
Chapman, Karen B.
Morin, Gregg B.
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Harley, Calvin B.
Harley, Calvin B.
Andrews, William H.
TITLE OF INVENTION: Human Telomerase Catalytic Subunit
NUMBER OF SEQUENCES: 727
CORRESPONDENCE ADDRESS:
ADDRESSE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
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OTHER INFORMATION: /note= "antisense hTRT molecule"
SEQUENCE DESCRIPTION: SEQ ID NO: 543:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 3.6%; Score 16.4; DB 1; Length 18; Best Local Similarity 94.4%; Pred. No. 1.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
NAME: Apple, Randolph Ted
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002610US
TELECOMMUNICATION INFORMATION:
                    APPLICATION NUMBER: US/09/432,503
FILING DATE: 02-Nov-1999
FILING DATE: 02-Nov-1999
FILING DATE: 02-Nov-1999
FILING DATE: 01-Nov-1999
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/851,843
FILING DATE: 06-MAY-1997
APPLICATION NUMBER: US 08/851,843
FILING DATE: 06-MAY-1997
APPLICATION NUMBER: US 08/851,050
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/911,312
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/912,951
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: WO PCT/US97/17618
FILING DATE: 01-OCT-1997
APPLICATION NUMBER: WO PCT/US97/17885
FILING DATE: 01-OCT-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Mismatches
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US-10-877-022-543/C
US-10-877-022-543, Application US/10877022
Publication No. US20040247613A1
GENERAL INFORMATION:
APPLICANT: Cech, Thomas R.
Lingner, Joachim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0200
INFORMATION FOR SEG ID NO: 543:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 149 CACCGTTCATTCTAGAGC 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 18 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                caccerrcarreragage 1
     PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-10-877-124-543
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TYPE: Nucleic Acid
STRANDEDNESS: Single
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COMPUTER READABLE FORM:
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                                       US-09-997-868-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-09-997-868-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
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Matches 1
            RESULT 226
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US-10-877-146-543
                                                                                                                                                                                                                                                                  CUDNIKE: USASA

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/877,146
FILING DATE: 24-UM-2004
PRIOR APPLICATION NUMBER: US/99/42,503
PRIOR APPLICATION NUMBER: US/99/44,419
FILING DATE: O-Nov-1999
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/844,419
FILING DATE: 16-APR-1997
APPLICATION NUMBER: US 08/854,050
FILING DATE: 10-MAY-1997
APPLICATION NUMBER: US 08/854,050
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/912,951
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/915,503
APPLICATION NUMBER: US 08/915,503
APPLICATION NUMBER: US 08/915,503
APPLICATION NUMBER: WO PCT/US97/17618
FILING DATE: 01-OCT-1997
APPLICATION NUMBER: WO PCT/US97/17618
FILING DATE: 01-OCT-1997
APPLICATION NUMBER: US 08/915,503
                           Andrews, William H.

PF INVENTION: Human Telomerase Catalytic Subunit
OF SEQUENCES: 727
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                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco STATE: California COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: Apple, Randolph Ted
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002610US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
Harley, Calvin B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   149 CACCGTTCATTCTAGAGC 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 18 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: 543:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: linear MOLECULE TYPE: DNA
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Best Local Similarity 94.49
Matches 17, Conservative
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Sequence 76, Application US/10071179

Sequence 76, Application US/20030108862A1

GENERAL INFORMATION:

GENERAL INFORMATION:

TITLE OF INVENTION: A NUCLEIC ACID ENCODING A RETINOBLASTOMA BINDING PROTEIN (RBP-7)

TITLE OF INVENTION: AND POLYMORPHIC MARKERS ASSOCIATED WITH SAID NUCLEIC ACID.

FILE REFERENCE: GENSET. 031A

CURRENT FILING DATE: 2002-02-07

PRIOR PAPLICATION NUMBER: BALLIER APPLICATION NUMBER: 09/345,882

PRIOR PAPLICATION NUMBER: EARLIER APPLICATION NUMBER: 05/091,315

PRIOR APPLICATION NUMBER: BALLIER APPLICATION NUMBER: US 60/091,315

PRIOR APPLICATION NUMBER: BALLIER APPLICATION NUMBER: US 60/111,909

PRIOR APPLICATION NUMBER: BALLIER PILING DATE: 1998-06-30

PRIOR APPLICATION NUMBER: BALLIER PILING DATE: 1998-12-10

NUMBER OF SEQ ID NOS: 140

SEQ ID NO 76

LENGTH: 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
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3.4%; Score 15.4; DB 1; Length 20;
Best Local Similarity 94.1%; Pred. No. 2.1e+02;
Matches 16; Conservative 0; Mismatches 1; Indels
                                                                                 Sequence 84, Application US/10714195
Publication No. US20050019785A1
GENERAL INFORMATION:
APPLICANT: Baker, Joffre
APPLICANT: Breer, Jose
TITLE OF INVENTION: GENE EXPRESSION PROFILLING OF EGFR
TITLE OF INVENTION: POSTITUE CANCER
TITLE OF INVENTION: POSTITUE CANCER
TITLE OF INVENTION: POSTITUE CANCER
TITLE OF INVENTION POSTITUE CANCER
TITLE OF INVENTION WINDER: US/10/714,195
CURRENT APPLICATION NUMBER: 60/427090
PRIOR APPLICATION NUMBER: 60/427090
PRIOR APPLICATION NUMBER: 60/427090
PRIOR FILING DATE: 2003-11-15
NUMBER OF SEQ ID NOS: 372
SEQ ID NO 84
LENGTH: 79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     166 CAAACAAAAATGTCAG 182
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity 54.2
Matches 32; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY: primer_bind LOCATION: 1..20
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US-10-071-179-76
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-10-714-195-84
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 71, Application US/10174020
Publication No. US20030232770A1
GENERAL INFORMATION:
APPLICANT: Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HYPOTHETICAL TUMOR ENDOTHELIAL MARKER EXF
TITLE REFERENCE: RTS-0369
CURRENT APPLICATION UNMBER: US/10/174,020
CURRENT FILING DATE: 2002-06-17
NUMBER OF SEQ ID NOS: 149
SEQ ID NO 71
LENGTH: 20
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Publication No. US20030232770A1
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Brett P. Monia
APPLICANT: Renneth W. Dobie
TITLE OF INVERTION: ANTISENSE MODULATION OF HYPOTHETICAL TUMOR ENDOTHELIAL MARKER EXF
FILE REFERENCE: RTS-0369
CURRENT APPLICATION NUMBER: US/10/174,020
CURRENT FILING DATE: 2002-06-17
NUMBER OF SEQ ID NOS: 149
SEQ ID NO 138
LENGTH: 20
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                                                                                                     OTHER INFORMATION: Description of Artificial Sequence: Synthesized OTHER INFORMATION: Amplification Primer Design based on EST Sequence; OTHER INFORMATION: GenBank Accession Number AA281296
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 15.8; DB 1; Length 20; Pred. No. 1.9e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; OTHER INFORMATION: Antisense Oligonucleotide US-10-174-020-71
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                431 CAGGACTCGGCTCACACATGC 451
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21 CAGGACTCGGCTCACCACTGC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 362 GGCCGCAGGAAGAGAACG 380
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   362 GGCCGCAGGAAGGAACG 380
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TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA ORGANISM: Artificial Sequence
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Best Local Similarity 89.5'
....hes 17; Conservative
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Best Local Similarity
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ORGANISM: H. sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-10-174-020-71/C
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                                                                          FEATURE:
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GENERAL INFORMATION:

APPLICANT: Ribozyme Pharmaceuticals, Inc.

APPLICANT: Ribozyme Pharmaceuticals, Inc.

APPLICANT: McSwiggen, James

APPLICANT: Fosnaugh, Kathy

TITLE OF INVENTION: Gene Expression Using Short Interfering RNA

TITLE OF INVENTION: Gene Expression Using Short Interfering RNA

CURRENT APPLICATION NUMBER: US/10/224,005

CURRENT APPLICATION NUMBER: US 60/315,315

PRIOR PILING DATE: 2002-08-20

PRIOR PILING DATE: 2001-08-28

SOFTWARE: PatentIn version 3.0

SEQ ID NO 315

LENGTH: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region US-10-224-005-315
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; Publication No. US20030170636A1
; GENERAL INFORMATION:
    APPLICANT: Susan M. Freier
    TITLE OF INVENTION: ANTISENSE MODULATION OF JAGGED 2 EXPRESSION
    TORRENT APPLICATION NUMBER: US/10/091,625
; CURRENT FILING DATE: 2002-03-05
; NUMBER OF SEQ ID NOS: 90
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: KOLler, Erich
APPLICANT: Shepard, Peter J.
TITLE OF INVENTION: JAGGED 2 INHIBITORS FOR INDUCING APOPTOSIS
FILE REFERENCE: ISPH-0660
CURRENT APPLICATION NUMBER: US/10/096,399A
NUMBER OF SEQ ID NOS: 91
SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 3.3%; Score 14.8; DB 1; Length 19; Best Local Similarity 88.9%; Pred. No. 2.2e+02; Matches 16; Conservative 0; Mismatches 2; Indels
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3.2%; Score 14.4; DB 1;
Best Local Similarity 93.8%; Pred. No. 1.7e+02;
Matches 15; Conservative 0; Mismatches 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21 GGGAGGGTGGTGGCCAT 38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19 GGGAGGGAGGTGGCCGT 2
          US20030143732A1
                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: RNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          265 CCCGGGGCTTCTCCGG 280
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; OTHER INFORMATION: PCR Primer US-10-091-625-4
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US-10-096-399A-4
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LENGTH: 16
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                                                                                                 TITLE OF INVENTION: A NUCLEIC ACID ENCODING A RETINOBLASTOMA BINDING PROTEIN (RBP-7) TITLE OF INVENTION: AND POLYMORPHIC MARKERS ASSOCIATED WITH SAID NUCLEIC ACID.
TITLE OF INVENTION: AND POLYMORPHIC MARKERS ASSOCIATED WITH SAID NUCLEIC ACID.
FILE REPERBNCE: 44.USS.DIV
CURRENT APPLICATION NUMBER: US/10/126,704
CURRENT FILING DATE: 1202-04-20
PRIOR APPLICATION NUMBER: US 60/091,315
PRIOR APPLICATION NUMBER: US 60/111,909
PRIOR PILING DATE: 1998-12-10
NUMBER OF SEQ ID NOS: 140
SOFTWARE: Patent.pm
SEQ ID NO 76
ENDERTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, James
APPLICANT: Posnaugh, Kathy
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Adenosine Al Receptor (AD
TITLE OF INVENTION: Gene Expression Using Short Interfering RNA
CURRENT APPLICATION NUMBER: US/10/224,005
RICHEAT FILING DATE: 2002-08-20
RICHEAT APPLICATION NUMBER: US 60/315,315
RICHEAT APPLICATION NUMBER: US 60/315,315
RICHEAT RILING DATE: 2001-08-28
NUMBER OF SEQ ID NOS: 347
SOFTWARE: PatentIn version 3.0
SEQ ID NO 154
LENGTH: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; LOCATION: 1..20
; OTHER INFORMATION: upstream amplification primer for SEQ 34, SEQ 55, SEQ 35, SEQ 56
US-10-126-704-76
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; OTHER INFORMATION: Description of Artificial Sequence: Target sequence/siNA sense
US-10-224-005-154
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3.4%; Score 15.4; DB 1; Length 20;
Best Local Similarity 94.1%; Pred. No. 2.1e+02;
Matches 16; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 19;
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3.3%; Score 14.8; DB 1; Length 1
Best Local Similarity 77.8%; Pred. No. 2.2e+02;
Matches 14; Conservative 2; Mismatches 2; Indels
WS-10-126-704-76
Sequence 76, Application US/10126704
Publication No. US20030170647A1
GENERAL INFORMATION:
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US-10-224-005-154
IS-10-224-005-154

; Sequence 154, Application US/10224005
; Publication No. US20030143732A1
; GENERAL INFORMATION:
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US-10-224-005-315/c
; Sequence 315, Application US/10224005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         166 CAAACAAAAATGTCAG 182
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 CAAACAATAAATGTCAG 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: RNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: primer_bind
LOCATION: 1..20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
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JERUSTAL INFORMATION
JERUSTAL INFORMATION
MONELL JOHN
APPLICANT: Cowsert, Lex M.
APPLICANT: Freier, Susan M.
APPLICANT: Samor, Henri M.
APPLICANT: Brooks, Douglas G.
APPLICANT: Brooks, Douglas G.
APPLICANT: Watt, Jacqueline R.
APPLICANT: Wickers, Timothy A.
TITLE OF INVENTION: DENTIFICATION OF GENETIC TARGETS FOR TITLE OF INVENTION: GENERATION OF OLIGONUCLECTIDES AND TITLE OF INVENTION: GENERATION OF OLIGONUCLECTIDES FOR GENE MODULATION FILE REPRENTED: 2003-03-12
CURRENT APPLICATION NUMBER: US/10/388,263
CURRENT APPLICATION NUMBER: US/10/388,263
CURRENT FILING DATE: 2003-03-12
NUMBER OF SEQ ID NOS: 947
SOFTWARE: FESTESEQ for Windows Version 4.0
SEQ ID NO 377
                                                                                                                                                                                                     Gaps
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                                                                                                                                                                                                                                                                                                                                                                                           Sequence 4. Application US/10461668
; Publication No. US20030207839A1
; BENERAL INFORMATION:
; APPLICANT: Suban M. Freier
; TITLE OF INVENTION: ANTISENSE MODULATION OF JAGGED 2 EXPRESSION
; FILE REFRENCE: RTS-0244
; CURRENT APLICATION NUMBER: US/10/461,668
; CURRENT FILING DATE: 2002-06-13
; PRIOR FILING DATE: 2002-03-05
; NUMBER OF SEQ ID NOS: 90
; SEQ ID NO 4
; LENGTH: 16
; TYPE: DNA
CREATISE ARTIFICIAL Sequence
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                                                                                                                                                   3.2%; Score 14.4; DB 1; Length 16; llarity 93.8%; Pred. No. 1.7e+02; Conservative 0; Mismatches 1; Indels
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3.2%; Score 14.4; DB 1;
Best Local Similarity 93.8%; Pred. No. 1.7e+02;
Matches 15; Conservative 0; Mismatches 1;
                                                           FEATURE:
; OTHER INFORMATION: Synthetic PCR primer
US-10-096-399A-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; Sequence 377, Application US/10388263; Publication No. US20030228597A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                  265 CCCGGGGCTTCTCCGG 280
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           265 CCCGGGCTTCTCCGG 280
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                     TYPE: DNA
ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; OTHER INFORMATION: PCR Primer US-10-461-668-4
                                                                                                                                                     Query Match
Best Local Similarity
Matches 15; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-10-388-263-377
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LENGTH: 16
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Sequence 145, Application US/09864785

CURRENT PRICE OF INVENTION: Exvels of NF-Kappa B

FILE REFERENCE: 400/022 (MBHB00-812-D)

CURRENT APPLICATION NUMBER: US/09/864,785

CURRENT APPLICATION NUMBER: 2001-05-23

NUMBER OF SEQ ID NOS: 3929

SOFTWARE: Patentin version 3.0

SEQ ID NO 145

LENGTH: 17
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Sequence 146, Application US/09664785

Sequence 146, Application US/09664785

PAPLICANT: Ribozyme Pharmaceuticals, Inc.

APPLICANT: Stinchcomb, Dan

APPLICANT: McSwiggen, Jim

TITLE OF INVENTION: Brymatic Nucleic Acid Treatment of Diseases or Conditions Relater

TITLE OF INVENTION: Brymatic Nucleic Acid Treatment of Diseases or Conditions Relater

TITLE OF INVENTION: Levels of NP-Kappa B

FILE REFERENCE: 400/022 (MBHB00-812-D)

CURRENT PRILIG DATE: 2001-05-23

NUMBER OF SEQ ID NOS: 3929

SOFTWARE PATENCE PATENCE: ACID NOS: 3929

SOFTWARE PATENCE PATENCE
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US-09-864-785-145
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; OTHER INFORMATION: Description of Artificial Sequence: Nucleic Acid
US-09-864-785-146
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                                                                                                                                                                      Query Match
3.2%; Score 14.4; DB 1; Length 16;
Best Local Similarity 93.8%; Pred. No. 1.7e+02;
Matches 15; Conservative 0; Mismatches 1; Indels
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; FEATURE:
; OTHER INFORMATION: PCR Primer
US-10-388-263-377
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ORGANISM: Artificial Sequence
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Best Local Similarity 75.0°
Matches 12; Conservative
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GENERAL INFORMATION

GENERAL INFORMATION

APPLICANT: Ribozyme Pharmaceuticals, Inc.

APPLICANT: Sandberg, Jennifer

APPLICANT: Gardon, Gilad

APPLICANT: Gordon, Gilad

APPLICANT: Gordon, Gilad

APPLICANT: Stinchomb, Dames

APPLICANT: Stinchomb, Dames

APPLICANT: Stinchomb, Dames

APPLICANT: Stinchomb, Dames

APPLICANT: McSwiggen, James

APPLICANT: WCSwiggen, James

PRIOR FILING DATE: 1995-10-26

PRIOR FILING DATE: 1995-08-10

PRIOR FILING DATE: 1999-08-10

PRIOR FILING DATE: 2001-05-29

PRIOR PRILING DATE: 2001-05-39

NUMBER OF SEQ ID NOS: 5999

SEQ ID NO 491

LENGTH: 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
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Publication No. US20030054365A1

GENURAL INFORMATION:

APPLICANT: AU, Minzhen

APPLICANT: AU, Minzhen

APPLICANT: Humphreys, Robert

TITLE OF INVENTION: ONDERR CELL VACCINE

FILE REFERENCE: U.S. Application 09/205, 995;

CURRENT APPLICATION NUMBER: US/10/054,387;

CURRENT APPLICATION NUMBER: 09/036,746

PRIOR APPLICATION NUMBER: 09/036,746

PRIOR PILING DATE: 1998-03-09

PRIOR FILING DATE: 1998-03-09

PRIOR FILING DATE: 1996-06-11

NUMBER OF SEQ ID NOS: 79

SOFTWARE: Patentin Ver. 2.0

SEQ ID NO 24
   US20040220128A1
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ORGANISM: Artificial Sequence
FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: RNA
ORGANISM: Homo Sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
Matches 15; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-10-712-633-491
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                                                                                                                                                                                                Sequence 7502, Application US/10138674

Sequence 7502, Application US/10138674

GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Roswiggen, Jim
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Jaime
APPLICANT: Stinchcomb, Jaime
APPLICANT: Stinchcomb, Jaime
APPLICANT: Stinchcomb, Jaime
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Jaime
APPLICANT: Stinchcomb, Jaime
APPLICANT: Stinchcomb, Jaime
APPLICANT: McSwiggen, Jim
APPLICANT: McSwiggen, Jim
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Jaim
APPLICANT: Stinchcomb, Jaime
TITLE OF INVENTION: McHod and Reagent for the Treatment of Diseases or Conditions ReFILE REFERENCE: MBHB00-876-N (400/049)
CURRENT APPLICATION NUMBER: US/10/138,674
NUMBER OF SEQ ID NOS: 20822
SOFTWARE: Patentin version 3.0
LENGTH: 17
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Sequence 7502, Application US/10287949A

Publication No. US20040102389A1

GENERAL INFORMATION:

APPLICANT: Ribozyme Pharmaceuticals, Inc.

APPLICANT: Ribozyme Pharmaceuticals, Inc.

APPLICANT: Bavco, Pam

APPLICANT: Binchcomb, Dan

APPLICANT: Stinchcomb, Dan

APPLICANT: Levels of Vascular Endothelial Growth Factor Receptor

TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor

CURRENT APPLICATION NUMBER: US/10/287,949A

NUMBER OF SEQ ID NOS: 20822

SOFTWARE: PatentIn version 3.0

SEQ ID NO 7502
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3.2%; Score 14.4; DB 1; Length 17;
Best Local Similarity 93.8%; Pred. No. 1.9e+02;
Matches 15; Conservative 0; Mismatches 1; Indels
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3.2%; Score 14.4; DB 1;
Best Local Similarity 93.8%; Pred. No. 1.9e+02;
Matches 15; Conservative 0; Mismatches 1;
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US-10-712-633-491/c
; Sequence 491, Application US/10712633
   131 CCTCGGCCTGCCGCCT 146
                                     28 GTGGTGGCCATTTTT 43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16 GTGATGGCCATTTTT 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: RNA
ORGANISM: Homo sapiens
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ORGANISM: Homo sapiens
                                                                                                                                                                                US-10-138-674-7502/c
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  Length 17;
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Edington, Brent E.
Mossingen, James A.
Merlo, Datricia Ann Owens
Guo, Lining
Skokut, Thomas A.
Young, Scott A.
Folkerts, Otto
Merlo, Donald J.
TITLE OF INVENTION: COMPOSITION AND METHODS FOR
MODULATION OF GENE EXPRESSION
IN PLANTS
NUMBER OF SEQUENCES: 1263
                                                  IndelB
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3.1%; Score 14; DB 1; Le ilarity 100.0%; Pred. No. 2.1e+02; Conservative 0; Mismatches 0;
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3.1%; Score 13.8; DB 1;
Best Local Similarity 64.7%; Pred. No. 2.2e+02;
Matches 11; Conservative 4; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM COMPATIBLE
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/961,077
FILING DATE: 21.569.2001
CLASSIFICATION: UNKNOWN.>
PRIOR APPLICATION NUMBER: 08/679,645
FILING DATE: July 12, 1996
APPLICATION NUMBER: 60/001,135
FILING DATE: July 12, 1996
APPLICATION NUMBER: 08/300,726
FILING DATE: September 2, 1994
ATTORNEY/AGENT INFORMATION:
NAME: WAZDLICY Richard J.
REGISTRATION NUMBER: 32,327
REERRENGE/DOCKET NUMBER: 32,327
TELECOMMUNICATION INFORMATION:
TELEPRANE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear SEQUENCE DESCRIPTION: SEQ ID NO: 826:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: Lyon & Lyon
STREET: 633 West Pifth Street
Suite 4700
                                                                                                                                                                                                                                        US-09-961-077-826
; Sequence 826, Application US/09961077
; Publication No. US20030014775A1
; GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO: 826:
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                                                                                               328 CTCTCGGGGGGGAG 341
                                                                                                                                             2 crcrceeeeccae 15
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Query Match
Best Local Similarity
Matches 14; Conserv
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US-10-093-958-22/C

Sequence 22, Application US/10093958

Sequence 22, Application US/10093958

Publication No. US20030044423A1

GENERAL INFORMATION:
APPLICANT: Glilies, Stephen
APPLICANT: Glilies, Stephen
APPLICANT: Glilies, Stephen
APPLICANT: Glilies, Stephen
TITLE OF INVENTION: Expression Technology for Proteins Containing a Hybrid Isotype Ar
TITLE OF INVENTION: Moiety
TITLE OF INVENTION: Moiety
TITLE OF INVENTION: WOMBER: US/10/093,958
CURRENT FILING DATE: 2002-03-07
PRIOR FILING DATE: 2001-03-07
NUMBER OF SEQ ID MOS: 50
NUMBER OF SEQ ID MOS: 50
SOFTWARE: Patentin version 3.0
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Publication No. US20050164243A1

GENERAL INFORMATION:

APPLICANT: Smith, Terry

APPLICANT: Mattin, Cara

APPLICANT: Mattin, Cara

APPLICANT: Mattin, Cara

APPLICANT: Wattin, Cara

APPLICANT: Wattin, Cara

APPLICANT: Van Der Weide, Marjo

TITLE OF INVENTION: Clinically important fungal pathogens

TITLE OF INVENTION: Clinically important fungal pathogens

FILE REFERENCE: 2551-49

CURRENT APPLICATION NUMBER: US/10/967,254

CURRENT APPLICATION NUMBER: US/09/662,462

PRIOR APPLICATION NUMBER: POP-19

PRIOR PILING DATE: 2000-09-15

PRIOR APPLICATION NUMBER: EP 99870109.8

PRIOR PILING DATE: 1099-05-28

PRIOR APPLICATION NUMBER: US 60/138,621

PRIOR APPLICATION NUMBER: US 60/138,621

PRIOR PILING DATE: 1999-06-11

NUMBER OF SEQ ID NOS: 43

SOFTWARE: PATEURIN Ver. 2.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; OTHER INFORMATION: forward primer for gamma 1 hinge region US-10-093-958-22
                           1; Indels
  93.8%; Pred. No. 2.1e+02; ive 0; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; TYPE: DNA
; ORGANISM: Aspergillus versicolor
US-10-967-254-21
                                                                           220 GGTCGCCTGCCCAGCC 235
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     300 GAAGAGTTGGGCTCTG 315
                                                                                                       1 GGTGGCCTGCCCAGCC 16
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  Best Local Similarity 93.8
Matches 15; Conservative
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LENGTH: 19
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RESULT 251
US-09-930-423-1471
US-09-930-423-1471

Sequence 1471, Application US/09930423

Publication No. US20030092003A1

GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Blatt, Larry
APPLICANT: McSwigen, Jim
TILLE OF INVENTION: Method and Reagent for the Treatment of Alzheimer's Disease
FILE REPERENCE: MBHB00, 918-A 400/027
CURRENT PELING DATE: 2001-08-15
NUMBER OF SEQ ID NOS: 4553

SOFTWARE: Patentin version 3.0

SEQ ID NO 1471
LENGTH: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 1472, Application US/09930423
; Sequence 1470, Application US/09930423
; Publication No. US20030092003A1
; GENERAL INFORMATION:
    APPLICANT: Ribozyme Pharmaceuticals, Inc.
    APPLICANT: Blatt, Larry
    APPLICANT: McSwiggen, Jim
    TITLE OF INVEXTURON: McHanco and Reagent for the Treatment of Alzheimer's Disease
    FILE REFERENCE: MRHB00, 918-A 400/027
    CURRENT APPLICATION NUMBER: US/09/930,423
    CURRENT RIING DATE: 2001-08-15
    NUMBER OF SEQ ID NOS: 4553
    SOFTWARE: PatentIn version 3.0
    SEQ ID NO 472
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                                                                                                                                                                                                                        3.1%; Score 13.8; DB 1; Length 17; 88.2%; Pred. No. 2.2e+02; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 3.1%; Score 13.8; DB 1; Length 17; Best Local Similarity 88.2%; Pred. No. 2.2e+02; Matches 15; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 17;
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88.2%; Pred. No. 2.2e+02;
cive 0; Mismatches 2;
              2001-08-15
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         CURRENT FILING DATE: 2001-08-1
NUMBER OF SEQ ID NOS: 4553
SOFTWARE: PatentIn version 3.0
SEQ ID NO 336
LENGTH: 17
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                                                                                                                                                                                                                                                                                                                                             17 GGCGGGCTGGGAGGGG 1
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                                                                                                                                                                                                                                         Best Local Similarity 88.2
Matches 15; Conservative
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Best Local Similarity 88.2°
Matches 15, Conservative
                                                                                                                         TYPE: RNA
CORGANISM: Homo Sapiene
US-09-930-423-336
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; ORGANISM: Homo Sapiens
US-09-930-423-1471
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ORGANISM: Homo Sapiens
US-09-930-423-1472
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US-09-930-423-1472
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                                                                                                           RESULT 248
US-09-848-754A-3581
S181, Application US/09848754A
Fublication No. US20030073207A1
GENERAL INFORMATION:
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
CURRENT APPLICATION NUMBER: US/09/848,754A
CURRENT FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 9645
SEQ ID NO 3581
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 5, Application US/09930423
Publication No. US200300920031
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Blat, Larry
APPLICANT: McSwiggen, Jim
FILE OF INVENTION: McHebod and Reagent for the Treatment of Alzheimer's Disease
FILE REFERENCE: MHBE00, 918-A 400/027
CURRENT APPLICATION NUMBER: US/09/930,423
CURRENT FILING DATE: 2001-08-15
NUMBER OF SEQ ID NOS: 4553
SOFTWARE: PatentIn version 3.0
SEQ ID NO 55
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Publication No. US20030092003A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Blatt, Larry
APPLICANT: MSWiggen, Jim
TITLE OF INVENTION: Method and Reagent for the Treatment of Alzheimer's Disease;
FILE REFERENCE: MBHB00,918-A 400/027
CURRENT APPLICATION NUMBER: US/09/930,423
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
3.1%; Score 13.8; DB 1;
Best Local Similarity 88.2%; Pred. No. 2.2e+02;
Matches 15; Conservative 0; Mismatches 2;
106 CGCTGACTTTCAGCGGG 122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         231 CAGCCCCCGAACCCCGC 247
                             1 CGCUGCCUUUCAGCUGG 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 CAGCCUCUGAACCCCGC 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: RNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: RNA
ORGANISM: Homo Sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity
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US-09-930-423-336/c
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Page 107

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APPLICANT: Ribozyme Pharmaceuticals Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relater
TITLE OF INVENTION: Hepatitis C Virus Infection
FILE REFERENCE: RPI 400/09/740,332
CURRENT APPLICATION NUMBER: US/09/740,332
CURRENT FILING DATE: 2001-03-26
NUMBER OF SEQ ID NOS: 9704
SOFTWARE: PatentIn version 3.0
SEQ ID NO 3130
LENGTH: 17
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Sequence 129, Application US/09792818

Sequence 129, Application Wo. US20030134806A1

GENERAL INFORMATION:

APPLICANT: Ribozyme Pharmaceuticals, Inc.

APPLICANT: Oon Carlowitz, Ira

APPLICANT: McWaggen, Jim

APPLICANT: Hamblin, Paul

APPLICANT: Bilis, Jonathan

ITLE OF INVENTION: Method and Reagent for the Inhibition of Grb-2-related with Insert

ITLE OF INVENTION: (GRID) Gene

CURRENT APPLICATION NUMBER: US/09/792,818

CURRENT FILING DATE: 2001-02-23

NUMBER OF SEQ ID NOS: 2304

SOFTWARE: Patentin version 3.0

IDEACH.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               3.1%; Score 13.8; DB 1; Length 17;
88.2%; Pred. No. 2.2e+02;
tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                        , OTHER INFORMATION: oligonucleotide substrate US-09-740-332-3130
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; Sequence 330, Application US/09792818; Publication No. US20030134806A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       201 CTCCCGGGGACCTGCGG 217
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20 TGGGAGGGGTGGTGGCC 36
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Von Carlowitz, Ira
McSwiggen, Jim
Hamblin, Paul
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                                                                                                                                                                                                                                                                  TYPE: RNA
ORGANISM: artificial sequence
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Best Local Similarity 88.2.
Then 15; Conservative
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Best Local Similarity 88.2.
Best Local 5; Conservative
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; ORGANISM: Homo sapiens
US-09-792-818-129
                                                                                                                                                                                                                                                                                                                                          NAME/KEY: misc_feature
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US-09-792-818-330/c
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APPLICANT:
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                                                                                    NGS-09-827-395A-630

Sequence 630, Application US/09827395A

Publication No. US20030113891A1

GENERAL INFORMATION:
APPLICANT: Lawrence Blatt
APPLICANT: James McSwiggen
APPLICANT: James McSwiggen
APPLICANT: James Chowriza
TITLE OF INVENTION: Method and Reagent for the Inhibition of NOGO and NOGO Receptor G
FILE REFERENCE: MBHB00-879-C (400/017)
CURRENT FILING DATE: 2001-04-05
PRIOR PLICATION NUMBER: 09/780,533
PRIOR PLLING DATE: 2001-04-05
PRIOR PLLING DATE: 2001-02-09
PRIOR PLLING DATE: 2001-02-09
PRIOR PLLICATION NUMBER: 60/181,797
PRIOR PLLING DATE: 2000-02-11
NUMBER OF SEQ ID NOS: 2617
SOFTWARE: Patentin version 3.0
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Publication No. US20030125270A1

GENERAL INFORMATION:

APPLICANT: Ribozyme Pharmaceuticals Inc.

TITLE OF INVENTION: Hepatitis C Virus Infection

CURRENT FILING DATE: 2001-03-26

NUMBER OF SEQ ID NOS: 9704

SEQ ID NO 800

LENGTH: 17
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ORGANISM: artificial sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: RNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: misc_feature LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity
Matches 13; Conserv
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US-09-740-332-3130/c
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242 CCCCGCCTGGAGGCCGC 258
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                                                               12 GGTGGGCCTGGGAGGGG
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Best Local Similarity 88.2
Matches 15; Conservative
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US-09-745-237A-1471
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; ORGANISM: Homo sapiens
US-09-745-237A-1472
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US-09-745-237A-1471
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LENGTH: 17
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Publication No. US20030143708A1

GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Blatt, Larry
APPLICANT: Blatt, Larry
APPLICANT: Blatt, Larry
APPLICANT: WcSwiggen, Jim
TITLE OF INVENTION: Method and Reagent for the Treatment of Alzheimer's Disease
CURRENT APPLICATION NUMBER: US/09/745,237A

CURRENT FILING DATE: 2002-04-15
SOPTWARE: PATENTING DATE: 2002-04-15
SOPTWARE: PATENTING DATE: 2002-04-15
SOPTWARE: PATENTING DATE: DOOR OF SEQUENTING DATE: DOOR DATE: 
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GENERAL INFORMATION.
APPLICANT: NIGORATION:
APPLICANT: Blatt, Larry
APPLICANT: Blatt, Larry
APPLICANT: Machine Control of Management for the Treatment of Alzheimer's Disease
TITLE OF INVENTION: Method 918-A)
CURRENT APPLICATION WUMBER: US/09/745,237A
NUMBER OF SEQ ID NOS: 4550
SOFTWARE FLING DATE: 2002-04-15
NUMBER OF SEQ ID NOS: 4550
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3.1%; Score 13.8; DB 1; Length 17;
Best Local Similarity 88.2%; Pred. No. 2.2e+02;
Matches 15; Conservative 0; Mismatches 2; Indels
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88.2%; Pred. No. 2.2e+02;
tive 0; Mismatches 2; Indels
                          FILE REFERENCE: MHHB00-901-A (400/013)
CURRENT APPLICATION NUMBER: US/09/792,818
CURRENT FILING DATE: 2001-02-23
NUMBER OF SEQ ID NOS: 2304
SOFTWARE: PatentIn version 3.0
TITLE OF INVENTION: (GRID) Gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      200 CCTCCCGGGGACCTGCG 216
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Best Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                                                                                                                                                         ; TYPE: RNA
; ORGANISM: Homo sapiens
US-09-792-818-330
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ORGANISM: Homo sapiens
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ORGANISM: Homo sapiens
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LENGTH: 17
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US-09-817-879-800
US-09-817-879-800
Sequence 800, Application US/09817879
FUBLICATION NO. US20030171311A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals Inc.
TITLE OF INVENTION: Hazymatic Nucleic Acid Treatment of Diseases or Conditions Related
TITLE OF INVENTION: Hepatitis C Virus Infection
FILE REFERENCE: MBHB00-801-F
Sequence 1471, Application US/09745237A
Sequence 1471, Application US/09745237A
Publication No. US20030143708A1
GENERAL INFORMATION:
APPLICANT: Blatt, Larry
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Method and Reagent for the Treatment of Alzheimer's Disease
CURRENT FILING MAPPLEO. 18709/745,237A
CURRENT FILING DATE: 2002-04-15
NUMBER OF SEQ ID NOS: 4550
SOFTWARE: Patentin version 3.0
SOFTWARE: Patentin version 3.0
SOFTWARE: Patentin version 3.0
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Bublication No. US20030143708A1

GENERAL INFORMATION:
APPLICANT: Blacty.
APPLICANT: Blatt, Larry
APPLICANT: Blatt, Larry
APPLICANT: Method and Reagent for the Treatment of Alzheimer's Disease
FILE REFERENCE: 400/007 (MRHB00-918-A)
CURRENT APPLICATION NUMBER: US/09/745,237A

CURRENT FILING DATE: 2002-04-15
NUMBER OF SEQ ID NOS: 4550
SOFTWARE PATENTIAN VESION 3.0
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Sequence 5872, Application US/10156306
| Sequence 5872, Application US/10156306
| Publication No. US20030119017A1
| Publication No. US20030119017A1
| Publication No. US20030119017A1
| APPLICANT: Ribozyme Pharmaceuticals, Inc.
| APPLICANT: McSwiggen, James
| TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relater TITLE OF INVENTION: Levels of IKK-Gamma and PKR | FILE REFERENCE: MBHB01-664-A (400/050) | CURRENT APPLICATION NUMBER: US/10/156,306 | UNMER OF SEQ ID NOS: 8013 | SOFTWARE: Patentin Version 3.0 | SOFTWARE: Patentin Version 3.0 | SEQ ID NO 5872
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LAPLICANT: Application US/1023006
Publication No. US2003019107741
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: METHOD AND REAGENT FOR THE TREATMENT OF ASTHMA AND ALLERGIC CONDIFILE REFERENCE: 400/056 (MBHB01-1110)
CURRENT APPLICATION NUMBER: US/10/230,006
CURRENT FILING DATE: 2002-11-18
PRIOR FILING DATE: 2001-08-28
NUMBER OF SEQ ID NOS: 2678
SOFTWARE: Patentin version 3.0
SSEQ ID NO 22177
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                                              Query Match 3.1%; Score 13.9; DB 1; Length 17; Best Local Similarity 76.5%; Pred. No. 2.2e+02; Matches 13; Conservative 2; Mismatches 2; Indels
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                                                                                                                                                   246 GCCTGGAGGCCGCGGTC 262
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Best Local Similarity 76.55
Matches 13, Conservative
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Best Local Similarity 88.2
Matches 15; Conservative
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; ORGANISM: Homo sapiens
US-10-156-306-5872
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: RNA
CRGANISM: Homo sapiens
US-10-230-006-2217
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US-10-230-006-2217/c
     US-10-156-306-5871
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US-10-156-306-5872
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Fublication No. US20030119017A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: MCSWiggen, James
TITLE OF INVENTION: Levels of IKK-Gamma and PKR
TITLE OF INVENTION: Levels of IKK-Gamma and PKR
FILE REFERENCE: MBHB01-664-A (400/050)
CURRENT APPLICATION NUMBER: US/10/156,306
CURRENT FILING DATE: 2002-05-28
NUMBER OF SEQ ID NOS: 8013
SOFTWARE: Patentin version 3.0
SEQ ID NO 5871
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Ribozyme Pharmaceuticals Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions FILLS PERERENCE: MBH800-801-F
CURRENT PRESENCE: MBH800-801-F
CURRENT FILING DATE: 2001-03-26
NUMBER OF SEQ ID NOS: 9703
SOFTWARE: Patentin version 3.0
ENGTHARE: Patentin version 3.0
ENGTHARE: 173
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                                                                                                                                                                                            FEATURE:
NAME/KEY: misc_feature
LOCATION:
PEATURE:
LOCATION:
OTHER INFORMATION: oligonucleotide substrate
US-09-817-879-800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: oligonucleotide substrate
CURRENT APPLICATION NUMBER: US/09/817,879
CURRENT FILING DATE: 2001-03-26
NUMBER OF SEQ ID NOS: 9703
SOFTWARE: Patentin version 3.0
LENGTH: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; Sequence 3130, Application US/09817879; Publication No. US20030171311A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                     432 AGGACTCGCCTCACACA 448
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: RNA
ORGANISM: artificial sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17 TGGGAGGGTGGTGGCC 1
                                                                                                                                                   TYPE: RNA ORGANISM: artificial sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: misc_feature LOCATION:
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ORGANISM: Homo sapiens
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US-10-156-306-5871
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US09436060A-16.rnpb.sl

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APPLICANT: Pamela, Pavco
APPLICANT: Partice, Lae
APPLICANT: Renneth, Draper
APPLICANT: Kenneth, Draper
APPLICANT: Elisabeth, Roberts
TITLE OF INVENTION: OLIGONUCLEOTIDE MEDIATED INHIBITION OF HEPATITIS B VIRUS AND HEPATITIE OF INVENTION: VIRUS REPLICATION
FILE REFERENCE: 400/042US (MBHEB02-249-E)
CURRENT APPLICATION NUMBER: US/10/669,841
CURRENT FILING DATE: 2003-09-23
PRIOR FILING DATE: 2003-01-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FRIOR FILING DATE: 2002-03-26

PRIOR FLING DATE: 2001-06-08

PRIOR FLING DATE: 2001-06-08

PRIOR APPLICATION NUMBER: US 60/296,876

PRIOR PELING DATE: 2001-06-08

PRIOR PELING DATE: 2001-10-24

PRIOR PELING DATE: 2001-10-24

PRIOR PELING DATE: 2001-12-02

PRIOR APPLICATION NUMBER: US 60/354,580

PRIOR APPLICATION NUMBER: US 60/363,124

PRIOR APPLICATION NUMBER: US 60/363,124

PRIOR PELING DATE: 2001-03-26

PRIOR PELING DATE: 2001-03-26

PRIOR PELING DATE: 2000-03-11

PRIOR PELING DATE: 2000-02-18

PRIOR PELING DATE: 2000-02-07

PRIOR PELING DATE: 2000-02-05

PRIOR PELING DATE: 2000-02-15

PRIOR PELING DATE: 2000-02-15
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OTHER INFORMATION: Description of Artificial Sequence: Nucleic Acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1; Mismatches
                                                                                                                                         Sequence 3399, Application US/10669841 Eublication No. US20040127446A1 GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc. APPLICANT: Lawrence, Blatt
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-10-669-841-5723/C
Sequence 5723, Application US/10669841
Sequence 5723, Application US/10669841
GENERAL INFORMATION:
APPLICANT: STATE Theoremetrics, Inc.
APPLICANT: Lawrence, Blatt
APPLICANT: Dennis, Macejak
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SEQ ID NO 3393
LENGTH: 17
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                  Dennis, Macejak
James, McSwiggen
David, Morrissey
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Best Local Similarity
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Matches
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                                                                   GENERAL INFORMATION:

APPLICANT: Hibozyme Pharmaceuticals, Inc.
APPLICANT: Hibozyme Blatt
APPLICANT: Lawrence Blatt
APPLICANT: Lawrence Blatt
APPLICANT: Dames McSwiggen
APPLICANT: Bharat Chowrira
CURRENT PILING DATE: 2001-05-06
PRIOR APPLICATION NUMBER: 09/827,395
PRIOR PILING DATE: 2001-02-09
PRIOR PILING DATE: 2001-02-09
PRIOR PILING DATE: 2001-02-09
PRIOR PILING DATE: 2001-02-09
PRIOR PILING DATE: 2000-02-01
PRIOR FILING DATE: 2000-03-01
NUMBER: PCT/US02/10512
NUMBER OF SEQ ID NOS: 2617
SOFTWARE: PatentIn version 3.0
FROM IN 630
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Sequence 2726, Application US/10712672

Rublication No. US20040102413A1

GENERAL INFORMATION:

APPLICANT: Ribozyme Pharmaceuticals, Inc.

APPLICANT: McSwiggen, Jim

APPLICANT: Chowrita, Bharat

APPLICANT: Stinchcomb, Dan

TITLE OF INVENTION: Method and Reagent for the Inhibition of Telomerase Enzyme

TILE REFERENCE: MBHB00-882-C (400/019)

CURRENT APPLICATION NUMBER: US/10/712,672

CURRENT APPLICATION NUMBER: US/10/712,672

FRIOR APPLICATION NUMBER: 60/197,769

PRIOR APPLICATION NUMBER: 60/197,769

PRIOR FILING DATE: 2000-08-31

PRIOR FILING DATE: 1999-08-31

NUMBER OF SEQ ID NOS: 5586

SOFWWARE: PatentIn version 3.0

LENGTH. 17
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3.1%; Score 13.8; DB 1; Length 17;
Best Local Similarity 88.2%; Pred. No. 2.2e+02;
Matches 15; Conservative 0; Mismatches 2; Indels
Sequence 630, Application US/10430882
Publication No. US20030203870A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           263 GGCCCGGGCCTTCTCCG 279
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; ORGANISM: Homo sapiens
US-10-430-882-630
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; ORGANISM: Homo sapiens
US-10-712-672-2726
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US-10-712-672-2726/c
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Sequence 27, Application US/10158160A
; Bublication No. US20030059805A1
; GENERAL INFORMATION:
    GENERAL INFORMATION:
    APPLICANT: RAPPOLD-HOERBRAND, GUDRUN
    APPLICANT: RAPPOLD-HOERBRAND, GUDRUN
    APPLICANT: RAPPOLD-HOERBRAND, GUDRUN
    TITLE OF INVENTION: HUMAN GROWTH GENE AND SHORT STATURE GENE REGION
    TITLE OF INVENTION: HUMAN GROWTH GENE AND SHORT STATURE GENE REGION
    TITLE OF INVENTION: HUMAN GROWTH GENE AND SHORT STATURE GENE REGION
    CURRENT APPLICATION NUMBER: 09/147,699
    PRIOR APPLICATION NUMBER: 09/147,699
    PRIOR APPLICATION NUMBER: 00/27,633
    PRIOR PILING DATE: 1997-09-29
    PRIOR PILING DATE: 1996-10-01
    PRIOR PILING DATE: 1996-10-01
    PRIOR PILING DATE: 1997-01-16
    NUMBER OF SEQ ID NOS: 55
    SEQ ID NO 27
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                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: WORD Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/961,077
FILING DATE: 21-569-2001
CLASSIFICATION: UNKNOWN:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/679,645
FILING DATE: July 12, 1996
APPLICATION NUMBER: 60/001,135
FILING DATE: July 13, 1995
APPLICATION NUMBER: 08/300,726
FILING DATE: SEPTEMBER: 08/300,726
                                                                                                                                                                                             COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 219/247
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; SEQUENCE DESCRIPTION: SEQ ID NO: 571:
US-09-961-077-571
                         ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 571:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 18 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   134 CGGCCTGCCGCCTTCCA 150
                                                                                                                                                                                                                                                                                                                                  storage
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CORRESPONDENCE ADDRESS
                                                                                                                                    CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity
Matches 14; Conservat
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                                                                                                           APPLICANT: Patrice, Lee
APPLICANT: Patrice, Lee
APPLICANT: Bairstee, Lee
APPLICANT: Elisabeth, Roberte
TITLE OF INVENTION: OLIGONUCLECTIDE MEDIATED INHIBITION OF HEPATITIS B VIRUS AND HEPA
TITLE OF INVENTION: OLIGONUCLECTION MEDIATED INHIBITION OF HEPATITIS B VIRUS REPLICATION WINBER: US/10/66,841
CURRENT APPLICATION NUMBER: US/10/66,841
PRIOR APPLICATION NUMBER: US/010/66,876
PRIOR PILING DATE: 2002-03-26
PRIOR PILING DATE: 2003-06-08
PRIOR PELING DATE: 2001-06-08
PRIOR PILING DATE: 2001-10-24
PRIOR PELING DATE: 2001-10-24
PRIOR APPLICATION NUMBER: US/0335,059
PRIOR PELING DATE: 2002-02-20
PRIOR APPLICATION NUMBER: US/0331,124
PRIOR APPLICATION NUMBER: US/0363,124
PRIOR PELING DATE: 2002-03-11
PRIOR APPLICATION NUMBER: US/0363,124
PRIOR PELING DATE: 2002-03-11
PRIOR PELING DATE: 2001-03-26
PRIOR PELING DATE: 2001-03-26
PRIOR PELING DATE: 2001-03-26
PRIOR PELING DATE: 2001-03-26
PRIOR PELING DATE: 2000-02-15
PRIOR PELING DATE: 2000-02-15
PRIOR PELING DATE: 2000-03-21
PRIOR PELING DATE: 2000-02-15
PRIOR PELING DATE: 2000-01-18
PRIOR PELING D
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    .
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: COMPOSITION AND METHODS FOR MODULATION OF GENE EXPRESSION IN PLANTS

NUMBER OF SEQUENCES: 1263
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 511, Application US/09961077
Publication No. US20030014775A1
GENERAL INFORMATION:
APPLICANT: Edington, Brent E.
MCSWiggen, James A.
MCSWiggen, James A.
Merlo, Patricia Ann Owens
Guo, Lining
Skokut, Thomas A.
Young, Scott A.
Folkerts, Otto
Merlo, Donald J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ), OTHER INFORMATION: oligonucleotide substrate US-10-669-841-5723
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20 TGGGAGGGGTGGTGGCC 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: RNA
ORGANISM: Artificial Sequence
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                                                                                                    Patrice, Lee
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: misc_feature
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US-09-740-332-3638/c

Sequence 3638, Application US/09740332

Sequence 3638, Application US/09740332

Sequence 3638, Application US/09740332

GENERAL INFORMATION:

APPLICANT: Ribozyme Pharmaceuticals Inc.

TITLE OF INVENTION: Hepatitis C Virus Infection

TITLE OF INVENTION: Hepatitis C Virus Infection

FILE REFERENCE: RP1 400/03

CURRENT APPLICATION NUMBER: US/09/740,332

CURRENT FILING DATE: 2001-03-26

NUMBER OF SEQ ID NOS: 9704

SOFTWARE: Patentin Version 3.0

TEMBER OF SEQ ID NOS: 9704

SOFTWARE: PATENTIN VERSION 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 276

US-09-792-818-429/C

US-09-792-818-429/C

Sequence 429, Application US/09792818

Sublication No. US20030134806A1

SUBJICANT: No. US20030134806A1

SEPLICANT: Jarvis, Thale

APPLICANT: Won Carlowitz, Ira

APPLICANT: Woskiggen, Jim

APPLICANT: Hamblin, Paul

ITILE OF INVENTION: Method and Reagent for the Inhibition of Grb-2-related with Insert

TILE OF INVENTION: Method and Reagent for the Inhibition of Grb-2-related with Insert

TILE OF INVENTION: MUBBR: US/09/792,818

CURRENT FILING DATE: 2001-02-23

NUMBER OF SEQ ID NOS: 2304

SOFTWARE: PatentIn version 3.0

SEQ ID NO 429

LENGTH: 17

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                              3.0%; Score 13.4; DB 1; Length 17; 80.0%; Pred. No. 2.4e+02; ive 2; Mismatches 1; Indels
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3.0%; Score 13.4; DB 1;
Best Local Similarity 93.3%; Pred. No. 2.4e+02;
Matches 14; Conservative 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; OTHER INFORMATION: oligonucleotide substrate US-09-740-332-3638
                                                                                                                                                        210 ACCTGCGGCGGGTCG 224
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: RNA
ORGANISM: artificial sequence
                                                                                                                                                                                                 2 ACCUGCGGCGGCUCG 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            210 ACCTGCGGCGGGTCG 224
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 132 CTCGGCCTGCCGCT 146
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                                                                                               Conservative
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ORGANISM: Homo sapiens
                    Query Match
Best Local Similarity
Matches 12; Conserv
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Sequence 1488, Application US/09864785

Sequence 1488, Application US/09864785

Sequence 1488, Application US/09864785

Sequence 1488, Application US/09864785

GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Brinchomb, Dan
APPLICANT: Draper, Ken
TITLE OF INVENTION: Levels of NF-Kappa B
TITLE OF INVENTION: Levels of NF-Kappa B
TITLE OF INVENTION: Levels of NF-Kappa B
CURRENT APPLICATION UNBER: US/09/864,785

CURRENT APPLICATION UNBER: US/09/864,785

CURRENT SEQ ID NOS: 3929

NUMBER OF SEQ ID NOS: 3929
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APPLICANT: Ribozyme Pharmaceuticals Inc.
APPLICANT: Ribozyme Pharmaceuticals Inc.
TITLE OF INVENTION: Braymatic Nucleic Acid Treatment of Diseases or Conditions Relate TITLE OF INVENTION: Hepatitis C Virus Infection
FILE REFERENCE: RPI 400/003
CURRENT APPLICATION NUMBER: US/09/740,332
CURRENT FILING DATE: 2001-03-26
NUMBER OF SEQ ID NOS: 9704
SOFTWARE: Patentin version 3.0
LENGTH: 17
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US-09-864-785-1488
                                                                                                                             ; OTHER INFORMATION: Description of Artificial Sequence: synthetic; CTHER INFORMATION: primer US-10-158-160A-27
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                                                                                                                                                                                                                                                                  3.1%; Score 13.8; DB 1; Length 18; 88.2%; Pred. No. 2.4e+02; tive 0; Mismatches 2; Indels
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3.0%; Score 13.4; DB 1; Length 17;
Best Local Similarity 73.3%; Pred. No. 2.48+02;
Matches 11; Conservative 3; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COTHER INFORMATION: oligonucleotide substrate US-09-740-332-917
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 274
US-09-740-332-917
Sequence 917, Application US/09740332
Publication No. US20030125270A1
                                                                                                                                                                                                                                                                                                                                                                                            332 CGGGGGGGGGGGGAGG 348
                                 TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              132 CTCGGCCTGCCGCT 146
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ORGANISM: artificial sequence
FEATURE:
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Best Local Similarity 88.2°
Matches 15; Conservative
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LENGTH: 17
LENGTH: 18
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| Publication No. US20030105314A1
| GENERAL INFORMATION |
| APPLICANT: Gaida, Marco |
| APPLICANT: Gaida, Marco |
| APPLICANT: Hall, Jeff |
| TITLE OF INVENTION: GENETIC TYPING OF THE HUMAN CYTOCHROME P450 2A6 GENE |
| TITLE OF INVENTION: AND RELATED MATERIALS AND METHODS |
| FILE REFERENCE: 4389-20 |
| CURRENT APPLICATION NUMBER: US/10/232,634 |
| CURRENT PILING DATE: 2002-08-30 |
| PRIOR PILICATION NUMBER: US/09/586,376 |
| NUMBER OF SEQ ID NOS: 29 |
| SOFTWARE: PatentIn Ver. 2.1
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Sequence 12, Application US/10232634

Publication No. US20030105314A1

GENERAL INFORMATION:

APPLICANT: Gaida, Marco

APPLICANT: Gaida, Marco

APPLICANT: Hall, Jeff

TITLE OF INVENTION: GENETIC TYPING OF THE HUMAN CYTOCHROME P450 2A6 GENE

TITLE OF INVENTION: AND RELATED MATERIALS AND METHODS

FILE REFERENCE: 4389-20

CURRENT FILING DATE: 2002-08-30

CURRENT FILING DATE: 2000-06-02

PRIOR FILING DATE: 2000-06-02
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
3.0%; Score 13.4; DB 1;
Best Local Similarity 93.3%; Pred. No. 2.4e+02;
Matches 14; Conservative 0; Mismatches 1;
TITLE OF INVENTION: Hepatitis C Virus Infection FILE REFERENCE: MBHB00-801-F CURRENT APPLICATION NUMBER: US/09/817,879 CURRENT FILING DATE: 2001-03-26 NUMBER OF SEQ ID NOS: 9703 SEQ ID NO 3638 LENGTH: 17
                                                                                                                                                                                                                                                                                                                                ; OTHER INFORMATION: oligonucleotide substrate US-09-817-879-3638
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3.0%;
                                                                                                                                                                                                                                ORGANISM: artificial sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            210 ACCTGCGGCGGGTCG 224
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SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14; Conservative
                                                                                                                                                                                                                                                       FEATURE:
NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
Matches 14: Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 280
US-10-232-634-11
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                                                                                                                                                                                                           TYPE: RNA
                                                                                                                                                                                                                                                                                                           LOCATION:
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                                                         Sequence 857, Application US/09792818
; Publication No. US2003134806A1
; GENERAL INFORMATION:
    APPLICANT: Ribozyme Pharmaceuticals, Inc.
; APPLICANT: Won Carlowitz, Ira
APPLICANT: Workwiggen, Jim
; APPLICANT: Hamblin, Paul
; APPLICANT: Hamblin, Paul
; APPLICANT: Hamblin, Paul
; APPLICANT: Hamblin, Paul
; TITLE OF INVENTION: Method and Reagent for the Inhibition of Grb-2-related with Inser
; TITLE OF INVENTION: (GRID) Gene
; TITLE OF INVENTION: (GRID) Gene
; FILE REFRENCE: MBHB00-901-A (400/013)
; CURRENT APPLICATION NUMBER: US/09/792,818
; CURRENT APPLICATION NUMBER: US/09/792,818
; SOFTWARE: Patentin version 3.0
; SEQ ID NO 857
; LENGTH: 17
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Subjection No. US20030171311A1

GENERAL INFORMATION:

APPLICANT: Ribozyme Pharmaceuticals Inc.

TITLE OF INVENTION: Hepatitis C Virus Infection

CURRENT PELLOATION WHOBER: US/09/817,879

CURRENT FILING DATE: 2001-03-26

NUMBER OF SEQ ID NOS: 9703

SEQ ID NO 917

LENGTH: 17
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; Publication No. US20030171311A1
; GENERAL INFORMATION:
; APPLICANT: Ribozyme Pharmaceuticals Inc.
; TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate
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3.0%; Score 13.4; DB 1;
Best Local Similarity 93.3%; Pred. No. 2.4e+02;
Matches 14; Conservative 0; Mismatches 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; OTHER INFORMATION: oligonucleotide substrate US-09-817-879-917
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       132 CTCGGCCTGCCGCCT 146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: artificial sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; TYPE: RNA
; ORGANISM: Homo sapiens
US-09-792-818-857
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: misc_feature LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-09-817-879-3638/c
                                                      US-09-792-818-857/c
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US-10-157-580A-68
Sequence 68, Application US/10157580A
Publication No. US20030124513A1
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: McSwiggen, Jim.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditing TITLE OF INVENTION: Related To Levels Of HIV
TITLE OF INVENTION: Related To Levels Of HIV
TITLE REFERENCE: MBH801-665-A (400/051)
CURRENT APPLICATION WUMBER: US/10/157,580A
NUMBER OF SEQ ID NOS: 170
SOFTWARE: Patentin version 3.0
SSEQ ID NO 68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 285

US-10-138-674-4559

Sequence 4559, Application US/10138674

Publication No. US2004007565A1

GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Rescoled, Dam
APPLICANT: Bacobed, Jam
APPLICANT: Escobed, Jam
APPLICANT: MSWITON: Levels of Vascular Endothelial Growth Factor Receptor
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHB00-876-N (400/049)
CURRENT FILING DATE: 2002-05-03

NUMBER OF SEQ ID NOS: 2082.

SEQ ID NO 94559

LENGTH: 17

LENGTH: 17
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Publication No. US20040077565A1

GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Roco, Pam
APPLICANT: Strickcomb, Dan
APPLICANT: Strickcomb, Dan
APPLICANT: Strokcomb, Jame
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Rel
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHB00-876-N (400/049)
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3.0%; Score 13.4; DB 1; Length 17;
Best Local Similarity 80.0%; Pred. No. 2.4e+02;
Matches 12; Conservative 2; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                        ; ORGANISM: Human immunodeficiency virus US-10-157-580A-68
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; ORGANISM: Homo sapiens
US-10-138-674-4559
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US-10-138-674-7501/c
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US-10-157-580A-38

15 Sequence 38, Application US/10157580A

1 Publication No. US2030124513A1

2 Sequence 38, Application US/10157580A

3 Publication No. US2030124513A1

3 GENERAL INFORMATION

4 APPLICANT: Ribozyme Pharmaceuticals, Inc.

5 APPLICANT: Ribozyme Pharmaceuticals, Inc.

6 TITLE OF INVENTION: Related To Levels Of HIV

7 TITLE OF INVENTION: Related To Levels Of HIV

7 TITLE OF INVENTION: Related To Levels Of HIV

7 TITLE OF INVENTION: Related To Levels Of HIV

7 TITLE OF INVENTION: Related To Levels Of HIV

7 TITLE OF SEQ ID NOSE: US/10/157,580A

7 CURRENT APPLICATION NUMBER: US/10/157,580A

7 SOFTWARE: PatentIn version 3.0

7 SEQ ID NO 38

7 FROTH: 17
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Sequence 27, Application US/10157580A
PSEQUENCE 27, Application US/10157580A
PUBLICANT: NO. US20030124513A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Related To Levels Of HIV
TITLE OF INVENTION: Related To Levels Of HIV
CURRENT APPLICATION NUMBER: US/10/157,580A
NUMBER OF SEQ ID NOS: 170
SOFTWARE: Patent In version 3.0
SSEQ ID NO 27
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3.0%; Score 13.4; DB 1; Length 17;
Best Local Similarity 80.0%; Pred. No. 2.4e+02;
Matches 12; Conservative 2; Mismatches 1; Indels
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; ORGANISM: Human immunodeficiency virus
US-10-157-580A-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            428 ACCCAGGACTCGGCT 442
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                                                                                                                                                                                                                                                      3 TTTGTCTCACCCTAA 17
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                                                                                                                 3.0%
Best Local Similarity 93.3%
Matches 14; Conservative
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Best Local Similarity 80.0°
Matches 12; Conservative
                  ; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-232-634-12
LENGTH: 17
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US09436060A-16.rnpb.sl

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APPLICANT: Kenneth, Debets
APPLICANT: Elisabeth, Roberts
TITLE OF INVENTION: OLIGONUCLECTIDE MEDIATED INHIBITION OF HEPATITIS B VIRUS AND HEPA
TITLE OF INVENTION: OLIGONUCLECTIDE MEDIATED INHIBITION OF HEPATITIS B VIRUS AND HEPA
FILE REFERENCE: 400/0420S (MRHB02-249-E)
CURRENT APPLICATION NUMBER: US/10/669, 441
CURRENT FILING DATE: 2003-09-23
FRIOR APPLICATION NUMBER: US 60/296, 876
FRIOR APPLICATION NUMBER: US 60/296, 876
FRIOR APPLICATION NUMBER: US 60/335, 059
FRIOR FILING DATE: 2001-10-60-8
FRIOR PELING DATE: 2001-10-60-8
FRIOR PELING DATE: 2002-03-10
FRIOR PELING DATE: 2000-13-26
FRIOR PELING DATE: 2000-13-26
FRIOR PELING DATE: 2000-13-26
FRIOR PELING DATE: 2000-13-26
FRIOR PELING DATE: 2000-03-16
FRIOR PELING DATE: 2000-03-16
FRIOR PELING DATE: 2000-03-16
FRIOR PELING DATE: 2000-02-15
FRIOR FILING D
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                                         Length 17;
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                                                                                                                        1; Indels
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                                         Score 13.4; DB 1;
Pred. No. 2.4e+02;
); Mismatches 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                  US-10-669-841-3510
Sequence 3510, Application US/10669841
Sequence 3510, Daplication US/10669841
GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Lawrence, Blatt
                                         Query Match 3.0%; Sco
Best Local Similarity 93.3%; Pro
Matches 14; Conservative 0;
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James, McSwiggen
David, Morrissey
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Patrice, Lee
Kenneth, Draper
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Best Local Similarity 80.0
Matches 12; Conservative
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US-10-287-949A-7501/C

Sequence 7501, Application US/10287949A

Publication No. US20040102389A1

GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Pavco, Pem Jim
APPLICANT: Stinchcomb, Dan
A
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Sequence 4559, Application US/10287949A

Publication No. US20040102389A1

GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Richcomb, Dan
APPLICANT: Bacobedo, Jaime
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHBDO-875-N (400/049)
CURRENT APPLICATION NUMBER: US/10/287,949A
CURRENT PILING DATE: 2003-04-11
NUMBER OF SEQ ID NOS: 20822

SOFTWARE: PatentIn version 3.0

SEQ ID NO 4559

LENGTH: 17
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                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 3.0%; Score 13.4; DB 1; Length 17; Best Local Similarity 93.3%; Pred. No. 2.4e+02; Matches 14; Conservative 0; Mismatches 1; Indels
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CURRENT APPLICATION NUMBER: US/10/138,674
CURRENT FILING DATE: 2002-05-03
NUMBER OF SEQ ID NOS: 20822
SOFTWARE: PatentIn version 3.0
SEQ ID NO 7501
LENGTH: 17
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Matches 13; Conservative
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US-10-138-674-7501
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ORGANISM: Homo sapiens
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-10-287-949A-4559
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TITLE OF INVENTION: NUCLEIC ACID BASED MODULATION OF VASCULAR ENDOTHELIAL GROWTH FACTY
TITLE OF INVENTION: RECEPTOR FOR THE TREATMENT OF ANGIOGENESIS RELATED DISEASES AND (
TITLE OF INVENTION: RECEPTOR FOR THE TREATMENT OF ANGIOGENESIS RELATED DISEASES AND (
CURRENT PEPLICATION NUMBER: US/10/712,633

CURRENT FILING DATE: 2003-11-13

PRIOR FILING DATE: 1995-10-26

PRIOR FILING DATE: 1995-01-08

PRIOR FILING DATE: 1995-01-08

PRIOR FILING DATE: 1995-01-08

PRIOR FILING DATE: 2000-11-07

PRIOR FILING DATE: 2001-05-29

PRIOR FILING DATE: 2001-05-29

PRIOR FILING DATE: 2001-05-29

PRIOR FILING DATE: 2001-11-30

PRIOR FILING DATE: 2001-05-29

PRIOR FILING DATE: 2001-11-30

PRIOR FILING DATE: 2001-05-29

PRIOR FILING DATE: 2001-05-29

PRIOR FILING DATE: 2001-05-29

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PRIOR FILING DATE: 2001-05-39

PRIOR PRIOR PRIOR PRIOR DATE: 2001-05-39

PRIOR PRIOR PRIOR DATE: 2001-05-39

PRIOR PRIOR PRIOR DATE: 2001-05-03

PRIOR PRIOR DATE: 2001-05-03

PRIOR PRIOR DATE: 2001-05-03
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Publication No. US200S0064495A1

GENERAL INFORMATION:

APPLICANT: Guida, Marco

FILLE OF INVENTION: GENETIC TYPING OF THE HUMAN CYTOCHROME P450 2A6

FILLE REFERENCE: DNA-20US2

CURRENT APPLICATION NUMBER: 10/20/32

PRIOR PLLING DATE: 2002-09-23

PRIOR PLLING DATE: 2002-08-30

PRIOR PLLING DATE: 2000-06-02

NUMBER OF SEQ ID NOS: 29

SOFTWARE: Patentin Ver. 2.1 and text editor

SEQ ID NO 11

LENGTH: 17
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Pred. No. 2.4e+02;
0; Mismatches 1; Indels
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US-10-949-004-12
: Sequence 12, Application US/10949004
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Best Local Similarity 93.3
Matches 14; Conservative
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US-10-712-633-490
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ORGANISM: Homo sapiens
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Matches 14; Conserv
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US-10-949-004-11
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                                                                                                                                                   APPLICANT: Lawrence, Blatt
APPLICANT: Lawrence, Blatt
APPLICANT: Lawrence, Blatt
APPLICANT: Lawrence, Blatt
APPLICANT: James, McSwiggen
APPLICANT: James, McSwiggen
APPLICANT: James, McSwiggen
APPLICANT: Pamela, Pavco
APPLICANT: Blisabeth, Roberts
TITLE OF INVENTION: OLIGONOLICEOTIDE WEDIATED INHIBITION OF HEPATITIS B VIRUS AND HEPA
TITLE OF INVENTION: OLIGONOLICEOTIDE WEDIATED INHIBITION OF HEPATITIS B VIRUS AND HEPA
TITLE REFERENCE: 400/0420S (MBHB02.249-8)
CURRENT APPLICATION NUMBER: PC7/0502/09187
PRIOR PELLING DATE: 2001-06-08
PRIOR PELLING DATE: 2001-10-04
PRIOR PELLING DATE: 2001-10-04
PRIOR PELLING DATE: 2001-12-05
PRIOR PELLING DATE: 2001-12-05
PRIOR PELLING DATE: 2001-12-05
PRIOR PELLING DATE: 2001-12-05
PRIOR PELLING DATE: 2000-02-22
PRIOR APPLICATION NUMBER: US 60/331,035
PRIOR PELLING DATE: 2000-02-12
PRIOR APPLICATION NUMBER: US 60/343,124
PRIOR PELLING DATE: 2000-01-21-31
PRIOR APPLICATION NUMBER: US 09/740,332
PRIOR PELLING DATE: 2000-01-21-38
PRIOR PELLING DATE: 2000-01-31
PRIOR APPLICATION NUMBER: US 09/504,321
PRIOR PELLING DATE: 2000-01-31
PRIOR PELLING DATE: 2000-01-31
PRIOR APPLICATION NUMBER: US 09/740,332
PRIOR PELLING DATE: 2000-01-31
PRIOR APPLICATION NUMBER: US 09/740,332
PRIOR PELLING DATE: 2000-01-31
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Publication No. US20040220128A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Pavco, Pamela
APPLICANT: Sandberg, Jennifer
APPLICANT: Gandberg, Jennifer
APPLICANT: Gordon, Gilad
APPLICANT: McSwiggen, James
                          Sequence 6231, Application US/10669841 Publication No. US20040127446A1 GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn version 3.0
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       210 ACCTGCGGCGGGTCG 224
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US-10-669-841-6231/c
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LENGTH: 17
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APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
TITLE OF INVENTION: Nacletc Acid Treatment of Diseases or Conditions Related to Level)
TITLE OF INVENTION: NAS, HERZ and HIV
FILE REPERENCE: 400/046-US (MHBB02-326-A)
CURRENT APPLICATION NUMBER: US/10/724,270
CURRENT PILING DATE: 2003-11-63
PRIOR FILING DATE: 2003-05-12
PRIOR FILING DATE: 2001-06-06
PRIOR PRIOR APPLICATION NUMBER: US 60/294,140
PRIOR FILING DATE: 2001-06-06
PRIOR FILING DATE: 2001-05-29
PRIOR FILING DATE: 2001-05-29
PRIOR FILING DATE: 2002-05-010
PRIOR PELICATION NUMBER: US 10/153,552
PRIOR PELICATION NUMBER: US 10/63,552
PRIOR PELICATION NUMBER: US 10/63,552
PRIOR PELICATION NUMBER: US 10/63,552
PRIOR PELICATION NUMBER: US 10/63,553
PRIOR PELICATION NUMBER: US 10/63,059
PRIOR PELICATION NUMBER: US 10/63
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APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, James
TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or Conditions Related to Level
TITLE OF INVENTION: RAS, HERZ and HIV
FILE REFERENCE: 400/046-US (MBHB02-326-A)
CURRENT APPLICATION NUMBER: US/10/724,270
CURRENT FILING DATE: 2003-11-26
PRIOR APPLICATION NUMBER: PCT/US02/16840
PRIOR FILING DATE: 2002-05-29
PRIOR FILING DATE: 2002-05-29
PRIOR FILING DATE: 2002-05-29
PRIOR FILING DATE: 2001-09-10
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   Mismatches
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                                                                              428 ACCCAGGACTCGGCT 442
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   Conservative
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US-10-724-270-6679
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Matches 12; Conserv
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US-10-724-270-6679
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   12;
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   Matches
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Sequence 6668, Application US/10724270

Publication No. US20050080031A1

GENERAL INFORMATION:

APPLICANT: SITRA THEADERLICE, Inc.

APPLICANT: SANSANGHON, James

ITILE OF INVENTION: NUCLEIC Acid Treatment of Diseases or Conditions Related to Level

TITLE OF INVENTION: NUMBER: US/10/724,270

CURRENT FILING DATE: 2003-11-26

FRIOR APPLICATION NUMBER: US/10/724,270

CURRENT FILING DATE: 2001-05-10

FRIOR APPLICATION NUMBER: US 60/239,440

FRIOR APPLICATION NUMBER: US 10/15,580

FRIOR APPLICATION NUMBER: US 10/15,580

FRIOR APPLICATION NUMBER: US 10/15,580

FRIOR FILING DATE: 2002-05-05

FRIOR APPLICATION NUMBER: US 10/15,580

FRIOR FILING DATE: 2003-05-05

FRIOR PLICATION NUMBER: US 10/15,580

FRIOR FILING DATE: 2003-05-05

FRIOR FILING DATE: 2003-06-05

FRIOR FILING DATE: 2003-06-10-3

                                                              APPLICANT: Guida, Marco
APPLICANT: Guida, Marco
TITLE OF INVENTION: GENETIC TYPING OF THE HUMAN CYTOCHROME P450 2A6 GENE
TITLE OF INVENTION: AND RELATED MATERIALS AND METHODS
FILE REPERENCE: DNA-20US2
CURRENT APPLICATION NUMBER: US/10/949,004
CURRENT FILING DATE: 2004-09-23
PRIOR FILING DATE: 2002-08-30
PRIOR PRIOR TOWN NUMBER: 09/586,376
PRIOR APPLICATION NUMBER: 09/586,376
PRIOR APPLICATION NUMBER: 09/586,376
PRIOR APPLICATION NUMBER: 09/586,376
PRIOR PAPLICATION NUMBER: 2000-06-02
NUMBER OF SEQ ID NOS: 29
SOFTWARE: PatentIN Ver. 2.1 and text editor
LUBRICH: 17
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Pred. No. 2.4e+02;
0; Mismatches 1; Indels
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Pred. No. 2.4e+02;
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Publication No. US20050064495A1 GENERAL INFORMATION:
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Best Local Similarity 93.3%;
Matches 14; Conservative
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; ORGANISM: Homo sapiens
US-10-724-270-6668
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
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GENERAL INFORMATION:
APPLICANT: BCKET, David J.
APPLICANT: BCKET, David J.
APPLICANT: BCKET, David J.
APPLICANT: Bnomett, C. Frank
APPLICANT: Bnomett, C. Frank
APPLICANT: Hanceak, Ronnie
APPLICANT: Hanceak, Ronnie
APPLICANT: Hanceak, Ronnie
APPLICANT: APPLICANT: Incothy
APPLICANT: Anderson, Kevin
ITLE OF INVENTION: Modulation Of Telomere Length By Oligonucleotides Having A G-Core
ITLE OF INVENTION: Modulation Of Telomere Length By Oligonucleotides
ITLE OF INVENTION: Modulation Of Telomere Length By Oligonucleotides
ITLE OF INVENTION: MOMBER: US/10/038,335
CURRENT FILING DATE: 1995-04-23
PRIOR APPLICATION NUMBER: 09/299,058
PRIOR PELING DATE: 1995-04-23
PRIOR PILING DATE: 1995-06-12
PRIOR PILING DATE: 1995-06-12
PRIOR PILING DATE: 1995-06-12
PRIOR PILING DATE: 1995-06-29
PRIOR PILING DATE: 1992-09-29
PRIOR PILING DATE: 1993-09-29
                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 13; DB 1; Length 13;
Pred. No. 1.5e+02;
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                               APPLICANT: Ahluwalia, Gurpreet S.
TITLE OF INVENTION: REDUCTION OF HAIR GROWTH
FILE REFERENCE: 00216-552001
CURRENT APPLICATION NUMBER: US/09/893,252
CURRENT FILING DATE: 2001-10-12
NUMBER OF SEQ ID NOS: 4
SOFTHARE: FastSEQ for Windows Version 4.0
SEQ ID NO 4
LENGTH: 13
                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 2.9%; Score 13; UB
Best Local Similarity 100.0%; Pred. No. 1.5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          46 CTAACCCTAACTG 58
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                                                                                                                                                                                                                                                                                                        TYPE: RNA
ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                    US-09-893-252-4
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Sequence 41, Application US/10911318

FOUNDication No. US20550130186A1

FOUNDication No. US20550130186A1

FOUNDICATION NO. US20550130186A1

FOUNDICANT: We Gene Technologies, Inc.

TITLE OF INVENTION: METHOD OF DETECTION CHIP AND FABRICATION METHOD THEREOF AND TITLE OF INVENTION: DETECTION OF DETECTION MENINGITIS AND PRIMER SET FOR MENINGITIS FILE REFERENCE: 1233-US-PA

CURRENT APPLICATION NUMBER: US/10/911,318

FRIOR FILING DATE: 2004-08-03

PRIOR PLING DATE: 2003-12-12

FRIOR PLING DATE: 2003-12-12

FRIOR PLING DATE: 2003-12-12
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PRIOR APPLICATION NUMBER: US 60/296,249
PRIOR FILING DATE: 2001-06-06
PRIOR PELING DATE: 2001-05-05
PRIOR PAPLICATION NUMBER: US 60/294,140
PRIOR PELING DATE: 2000-05-29
PRIOR FILING DATE: 2002-09-10
PRIOR PAPLICATION NUMBER: US 10/163,552
PRIOR PELING DATE: 2002-06-06
PRIOR PILING DATE: 2002-05-29
PRIOR PILING DATE: 2002-05-29
PRIOR PILING DATE: 2002-05-29
PRIOR PILING DATE: 2002-10-23
PRIOR APPLICATION NUMBER: US 10/444,853
PRIOR APPLICATION NUMBER: US 10/444,853
PRIOR PILING DATE: 2003-05-23
PRIOR PILING DATE: 2003-05-23
PRIOR PILING DATE: 2003-04-16
PRIOR PILING DATE: 2003-04-16
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 6810
COP THE NO. 2003-05-10
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US-09-893-252-4/c
i Sequence 4, Application US/09893252;
Publication No. US20030012755A1
; GENERAL INFORMATION:
; APPLICANT: Styczynski, Peter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               428 ACCCAGGACTCGGCT 442
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SOFTWARE: Patentin version 3.3
SEQ ID NO 41
LENGTH: 17
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Matches 12; Conservative
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US-10-724-270-6709
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US-10-463-076-2/C

US-10-463-076-2/C

Sequence 2, Application US/10463076

Publication NO. 220030212032A1

GENERAL INFORMATION:

APPLICANT: Geron Corporation

APPLICANT: Gryaznov, Sergei

APPLICANT: Pongracz, Krisztina

APPLICANT: Matray, Tracey

ITILE OF INVENTION: Oligonucleotide N3'-->PS' Thiophosphoramidates: Their Synthesis a

TITLE REPERBENCE: 039/004C

CURRENT APPLICATION NUMBER: US/10/463,076

CURRENT APPLICATION NUMBER: US 60/153,201

PRIOR FILING DATE: 1999-09-10

PRIOR PELING DATE: 1999-09-10

PRIOR PELING DATE: 1999-09-10

PRIOR APPLICATION NUMBER: US 60/153,201

PRIOR APPLICATION NUMBER: US 60/153,201

PRIOR SEQ ID NOS: 9

NUMBER OF SEQ ID NOS: 9

SOFTWARE: Patentin version 3.1
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; Publication No. US20030212032A1
; Sequence 8, Application US/10463076
; Publication No. US20030212032A1
; GENERAL INPORMATION:
; APPLICANT: Geron Corporation
; APPLICANT: Gryaznov, Sergei
; APPLICANT: Bongracz, Krisztina
; APPLICANT: PONGRACZ, Krisztina
; APPLICANT: MARIZAY, Tracey
; TILLE OF INVENTION: Oligonucleotide N3'-->P5' Thiophosphoramidates: Their Synthesis ;
; CURRENT APPLICATION NUMBER: US 00/6-17
; PRIOR FILING DATE: 2003-06-17
; PRIOR FILING DATE: 1999-09-10
; PRIOR FILING DATE: 1999-09-10
; PRIOR FILING DATE: 1999-10-19
; PRIOR FILING DATE: 1999-10-19
; NUMBER OF SEQ ID NOS: 9
; SOFTWARE: Patentin version 3.1
; SEQ ID NO 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; OTHER INFORMATION: Synthetic oligonucleotide with potential inhibition activity US-10-463-076-2
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ORGANISM: Artificial Sequence
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13 TTGTCTAACCCTA 1
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Best Local Similarity
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             APPLICANT: Brown-bried, Nonnie
APPLICANT: Brown-bried, Nonnie
APPLICANT: Hanecak, Ronnie
APPLICANT: Hanecak, Ronnie
APPLICANT: Hanecak, Ronnie
APPLICANT: Chiang, Ming-yi
APPLICANT: Chiang, Ming-yi
APPLICANT: Anderson, Kevin
TITLE OF INVENTION: Modulation Of Telomere Length By Oligonuclectides Having A G-Core
FILE RERERENCE: 1518-4976
FILE REPERENCE: 1518-4976
CURRENT APPLICATION NUMBER: 09/299,058
PRIOR APPLICATION NUMBER: 09/299,058
PRIOR PALICATION NUMBER: 08/403,888
PRIOR PPLING DATE: 1995-04-23
PRIOR PPLING DATE: 1993-09-29
PRIOR PLING DATE: 1993-09-29
PRIOR PPLING DATE: 1993-09-29
PRIOR PPLING DATE: 1992-09-29
PRIOR PLING DATE: 1992-09-29
NUMBER: OF SEQ ID NOS: 10
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US-10-25-35-3/C
Sequence 3, Application US/1025535
Fublication No. US2030138814A1
GENERAL INFORMATION:
APPLICANT: Geron Corporation
APPLICANT: Granov, Serget
APPLICANT: Tolman, Richard L.
APPLICANT: Morin, Gregg B.
TITLE OF INVENITION: Oligonucleotide Conjugates
FILE REFERENCE: 072/002P
CURRENT APPLICATION NUMBER: US/10/255,535
CURRENT APPLICATION NUMBER: S002-09-25
PRIOR FILING DATE: 2002-09-25
PRIOR FILING DATE: 2002-09-21
PRIOR FILING DATE: 2002-09-23
NUMBER OF SEQ ID NOS: 19
SOFTWARE: PLING DATE: 2001-03-23
NUMBER OF SEQ ID NOS: 19
SEQ ID NO 3
LENGTH: 13
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     Wyatt, Jacqueline
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LENGTH: 13
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Separation APPLICANT: Geron Corporation
APPLICANT: Gryaznov, Sergei
APPLICANT: Gryaznov, Sergei
APPLICANT: Gryaznov, Sergei
APPLICANT: Gryaznov, Krisztina
APPLICANT: Matray, Tracey
TITLE OF INVENTION: Oligonucleotide N3'-->P5' Thiophosphoramidates: Their Synthesis as
TITLE OF INVENTION: Oligonucleotide N3'-->P5' Thiophosphoramidates: Their Synthesis as
TITLE OF INVENTION: Oligonucleotide N3'-->P5' Thiophosphoramidates: Their Synthesis as
FILE REPRENCE: 030-06-17
FILE REPRENCE: 030-06-17
FRIOR APPLICATION NUMBER: US 09/657,445
FRIOR PILING DATE: 1099-09-10
FRIOR APPLICATION NUMBER: US 60/153,201
FRIOR APPLICATION NUMBER: US 60/160,444
FRIOR FILING DATE: 1999-10-19
NUMBER OF SEO ID NOS: 9
SOFTWARE: Patentin version 3.1
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GENERAL INFORMATION NO. US20050049408A1

APPLICANT: Geron Corporation

APPLICANT: Graznov, Sergei

APPLICANT: Matray, Tracey

TTLE OF INVENTION: Oligonuclectide N3'-->P5' Thiophosphoramidates: Their Synthesis and

TTLE OF INVENTION: Oligonuclectide N3'-->P5' Thiophosphoramidates: Their Synthesis and

FILE REPERBORCE: 039/005C

CURRENT APPLICATION NUMBER: US/10/967,755

CURRENT APPLICATION NUMBER: US 09/657,445

FRIOR PELING DATE: 2003-06-17

FRIOR PELING DATE: 1999-09-10

FRIOR PELING DATE: 1999-09-10

FRIOR APPLICATION NUMBER: US 60/153,201

FRIOR PILING DATE: 1999-10-19

FRIOR FILING DATE: 1999-10-19

FRIOR FILING DATE: 1999-10-19

FRIOR FILING DATE: 1999-10-19

SEQ ID NOS: 9

SOFTWARE: Patentin version 3.1
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100.0%; Pred. No. 1.5e+02;
Live 0; Mismatches 0; Indels
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Best Local Similarity 100.
Matches 13; Conservative
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Matches 13; Conservative
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LENGTH: 13
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                                                                                                                                                                                                                        Sequence 18, Application US/10181823

Publication No. US20040126752A1

GENERAL INRORMATION:

APPLICANT: GEYAZNOV, Sergei

APPLICANT: GEYAZNOV, Sergei

TITLE OF INVENTION: 2'-Arabino-Fluoroligonucleotide N3'-->P5' Phosphoramidates: Thei

TITLE OF INVENTION: Synthesis and Use

FILE REFERENCE: 049/002

CURRENT APPLICATION NUMBER: V3/10/181,823

CURRENT FILING DATE: 2003-12-29

PRIOR APPLICATION NUMBER: PCT/US01/01918

PRIOR FILING DATE: 2001-01-19

NUMBER: PEQ ID NOS: 23

SOFTWARE: PatentIn version 3.1

SEQ ID NO 18
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| Bublication No. US20040126752A1
| Publication No. US20040126752A1
| Publication No. US20040126752A1
| Publication No. US20040126752A1
| APPLICANT: Grown Corporation
| TITLE OF INVENTION: 2'-Arabino-Fluoroligonucleotide N3'-->P5' Phosphoramidates: Their
| TITLE OF INVENTION: Synthesis and Use
| TITLE OF INVENTION: Synthesis and
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Best Local Similarity 100.0%; Pred. No. 1.5e+02;
Matches 13; Conservative 0; Mismatches 0;
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2.9%; Score 13; DB
Best Local Similarity 100.0%; Pred. No. 1.5
Matches 13; Conservative 0; Mismatches
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   46 CTAACCCTAACTG 58
                                          13 CTAACCCTAACTG 1
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                                                                                                                                                                                                      US-10-181-823-18/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 306
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of Diseases or Conditions Related to Level
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Sequence 2188, Application US/10724270

Publication No. US20050080031A1

GENERAL INCORPATION:

APPLICANT: SITHA THE APPLICANT: GITHA THE APPLICANT:

APPLICANT: MCSAIGGRAN, James

TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or Conditions Related to Level

TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or Conditions Related to Level

TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or Conditions Related to Level

TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or Conditions Related to Level

TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or Conditions Related to Level

FILE REFERENCE: 400/046-US (MRH802-226-A)

CURRENT PAPLICATION NUMBER: US 60/296, 249

PRIOR FILING DATE: 2001-05-05

PRIOR APPLICATION NUMBER: US 60/294, 140

PRIOR APPLICATION NUMBER: US 10/238, 700

PRIOR FILING DATE: 2002-09-10

PRIOR FILING DATE: 2003-09-10

PRIOR PRIOR APPLICATION NUMBER: US 10/157, 580

PRIOR PRIOR APPLICATION NUMBER: US 10/14, 853

PRIOR PRIING DATE: 2003-09-23

PRIOR APPLICATION NUMBER: US 10/14, 853

PRIOR PRIOR APPLICATION NUMBER: US 10/417, 012

PRIOR APPLICATION NUMBER: US 10/417, 012

PRIOR APPLICATION NUMBER: US 10/416, 853

PRIOR APPLICATION NUMBER: US 10/416, 853

PRIOR APPLICATION NUMBER: US 10/416, 861

PRIOR APPLICATION NUMBER: US 10/416, 861

PRIOR PRIOR APPLICATION NUMBER: US 10/416, 861

PRIOR PRI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1; Le
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100.0%; Pred. No. 2.6e+02;
tive 0; Mismatches 0;
                     TITLE OF INVENTION: Vames
TITLE OF INVENTION: Nucleic Acid Treatment of
FILE REFRENCE: 400/057 (MBHB01-1158-A)
CURRENT APPLICATION NUMBER: US/10/238,700
CURRENT FILING DATE: 2002-09-18
PRIOR FILING DATE: 2002-05-29
PRIOR FILING DATE: 2001-09-10
PRIOR FILING DATE: 2001-09-10
PRIOR FILING DATE: 2001-09-10
SOFTWARE: Patentin version 3.0
SEQ ID NO 3509
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2.9%; Score 13; DB 100.0%; Pred. No. 2.6 iive 0; Mismatches
      APPLICANT: Ribozyme Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   225 CCTGCCCAGCCCC 237
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Best Local Similarity 100.C
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 100.
Matches 13, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        13 cereceasecee 1
                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: RNA
CAGANISM: Homo sapiens
US-10-238-700-3509
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                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: A POLYNUCLEIC ACID ISOLATED FROM A
PORCINE REPRODUCTIVE AND RESPIRATORY SYNDROME VIRUS (PRRSV)
A PROTEIN ENCODED BY THE POLYNUCLEIC ACID, A VACCINE
PREPARED FROM OR CONTAINING THE POLYNUCLEIC ACID OR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 77
CORRESPONDENCE ADDRESS:
ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
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STREET: 1755 S. Jefferson Davis Highway, Suite 400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
CORPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION NUMBER: US/10/618,779
FILING DATE: 15-Jul-2003
CLASSIFICATION NUMBER: US/08/301,435
FILING DATE: 01-SEP-1994
APPLICATION NUMBER: US/08/31,625
FILING DATE: 01-SEP-1993
ATTORNEY/AGENT INFORMATION:
NAMEN CONTINUED: US 08/131,625
FILING DATE: 05-OCT-1993
ATTORNEY/AGENT INFORMATION:
NAMEN CONTINUED: US 08/131,625
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 2.9%; Score 13; DB 1; Length 16; Best Local Similarity 100.0%; Pred. No. 2.3e+02; Matches 13; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Lavalleye, Jean-Paul M.P.
REGIGYRATION NUMBER: 31,451
REFERENCE/DOCKET NUMBER: 4625-021-55X CIP
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
SEQUENCE DESCRIPTION: SEQ ID NO: 34:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 309
US-10-238-700-3509/c
; Sequence 3509, Application US/10238700
; Publication No. US20030153521A1
                                                                                                                                                                                                                                                                                                    MENG, XIANG-JIN
HALBUR, PATRICK G.
MOROZOV, IGOR
LUM, MELISSA A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (703) 413-3000
                                                                                                                                                                              Sequence 34, Application US/10618779
Publication No. US20050175633A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (703) 413-2220
TELEX: 248855 OPAT UR
INFORMATION FOR SEQ ID NO: 34:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 16 base pairs
TYPE: nucleic acid
STRANDEDNESS: unknown
                                                                                                                                                                                                                                                                     APPLICANT: PAUL, PREM S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             268 GGGCTTCTCCGG 280
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4 GGGGCTTCTCCGG 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: Arlington
STATE: Virginia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: U.S.A.
13 CTAACCCTAACTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEPHONE:
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                                                                                                                      RESULT 308
US-10-618-779-34
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APPLICANT: StinchComb, Dan
TITLE OF INVENTION: Method and Reagent for the Inhibition of Telomerase Enzyme
TITLE OF INVENTION: Method and Reagent for the Inhibition of Telomerase Enzyme
TITLE OF INVENTION: Method and Reagent for the Inhibition of Telomerase Enzyme
CURRENT APPLICATION NUMBER: US/10/712,672
CURRENT PILING DATE: 2003-11-13
PRIOR PAPLICATION NUMBER: 06/197,769
PRIOR FILING DATE: 2000-04-14
PRIOR FILING DATE: 1999-08-31
NUMBER OF SEQ ID NOS: 5586
SOFTWARE: PATCHING VERSION 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 61, Application US/10483958

Sequence 61, Application US/10483958

Publication No. US2004025436341

SERNERAL INFORMATION:

APPLICANT: PRICE FOUNDATION LIMITED

APPLICANT: PRACER, Mercadith

APPLICANT: BERGEN, Andrew W.

TITLE OF INVENTION: GENES AND SNPS ASSOCIATED WITH EATING DISORDERS

FILE REPERENCE: 33061-5005-US

CURRENT FILING DATE: 2004-01-16

PRIOR APPLICATION NUMBER: US/10/483,958

CURRENT FILING DATE: 2001-07-16

PRIOR PLING DATE: 2001-07-16

PRIOR APPLICATION NUMBER: US 60/305,153

PRIOR PELING DATE: 2001-07-16

PRIOR PELING DATE: 2001-07-16

PRIOR FILING DATE: 2001-10-13

PRIOR FILING DATE: 2001-11-19

PRIOR FILING DATE: 2001-11-19

PRIOR FILING DATE: 2001-11-19

PRIOR FILING DATE: 2001-11-19

PRIOR FILING DATE: 2001-12-19

PRIOR FILING DATE: 2001-12-19

PRIOR FILING DATE: 2001-12-19

PRIOR FILING DATE: 2001-12-19

SOFTWARE: PALENTIN VUMBER: US 60/340,844

PRIOR FILING DATE: 2001-12-19

SOFTWARE: PALENTIN VUMBER: US 60/340,844

PRIOR FILING DATE: 2001-12-19

BRIOR FILING DATE: 2001-12-19

SEQ ID NO 61

BRIGHTH 16

FUNDER PRIOR FILING DATE: 2001-12-19

SEQ ID NO 61

BRIGHTH 16

FUNDER PRIOR FILING DATE: 2001-12-19

BRIGHTH EATING DATE: 2001-12-19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2.8%; Score 12.8; DB 1; Length 16; ilarity 87.5%; Pred. No. 2.4e+02; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; OTHER INFORMATION: HTR1D probe: FAM and MGB tagged US-10-483-958-61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              247 CCTGGAGGCCGCGGTC 262
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA ORGANISM: Artificial sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2 GGTTGCGGAGGGTGGG 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: RNA
GRGANISM: Homo sapiens
US-10-712-672-1787
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 14; Conserval
                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQ ID NO 1787
LENGTH: 16
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81.2%; Pred. No. 2.4e+02;
iive 1; Mismatches 2; Indels
                              WESULT 311
US-08-887-505-159
Sequence 159, Application US/08887505
Fublication No. US20020081577A1
GENERAL INFORMATION:
APPLICANT: Kilkuskie, Robert E.
APPLICANT: Frank, Bruce L.
APPLICANT: Goodchild, John
APPLICANT: Roberts, Peter C.
APPLICANT: Roberts, Peter C.
APPLICANT: Roberts, Poto. US20020081577A11 A.
APPLICANT: ROBERTS. POTOR OLIGONUCLEOTIDES SPECIFIC FOR TITLE OF INVENTION: OLIGONUCLEOTIDES SPECIFIC FOR NUMBER OF SEQUENCES: 172
CORRESPONDENCE ADDRESS:
ADDRESSEE: Hale and DOTT LLP
STREET: 60 State Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/887,505
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE:

CLASSIFICATION: 514

PRIOR APPLICATION: 514

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/471,968

FILING DATE: 06-JUN-1995

ATTONNEY/AGENT INFORMATION:

NAME: KAETHER, AND-LOUISE

REGISTRATION NUMBER: 33,523

REFERENCE/DOCKET NUMBER: HYZ-040CIP

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:

TELEFAX: (617) 526-600

TELEFAX: (617) 526-500

INFORMATION FOR SEQ ID NO: 159:

SEQUENCE CHARACTERISTICS:

LENGTH: 16 base pairs

TYPE: mucleic acid

TYPE: mucleic acid
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Publication No. US20040102413A1
GENERAL INFORMATION:
APPLICANT: Riboxyme Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        199 CCCTCCCGGGGACCTG 214
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13 CCTGCCCAGCCCC 1
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Best Local Similarity 81.2
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 02109
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy of
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STATE: MA
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-10-712-672-1787
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US-08-887-505-159
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RESULT 314

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VESULY 316

US-09-780-533A-27/C

US-09-780-533A-27/C

Sequence 27, Application US/09780533A

Publication No. US2000066611A1

GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Blact, Larry
APPLICANT: McSwiggen, Jim
APPLICANT: Wethod and Reagent for the Inhibition of NOGO Gene
FILE REFERENCE: MBH800,878-A (400/011)
CURRENT APPLICATION NUMBER: US 60/181,797
PRIOR PILING DATE: 2001-02-09
PRIOR PILING DATE: 2000-02-11
NUMBER OF SEQ ID NOS: 6679
SOFTWARE: PatentIn version 3.0
SEQ ID NO 27
LENGTH: 17
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Pred. No. 2.7e+02;
0; Mismatches 2;
CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/961,077
FILING DATE: 21-8ep-2001
CLASSIFICATION: 4Uhrnown>
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 08/679,645
FILING DATE: July 12, 1996
APPLICATION NUMBER: 60/001,135
FILING DATE: July 13, 1995
APPLICATION NUMBER: 08/300,726
FILING DATE: September 2, 1994
ATTORNEY/AGENT INFORMATION:
NAME: WATBURY, RICHARD 3.
REGISTRATION NUMBER: 32,327
REGISTRATION NUMBER: 21,327
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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US-09-961-077-828
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; Sequence 58, Application US/09780533A
                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 828:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 17 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  107 GCTGACTTTCAGCGGG 122
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Best Local Similarity 87.5
Matches 14; Conservative
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US-09-780-533A-27
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Best Local Similarity
Matches 10; Conserva
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2.8%; Score 12.8; DB 1; Length 16;
Best Local Similarity 87.5%; Pred. No. 2.4e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
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Merlo, Donald J.
TITLE OF INVENTION: COMPOSITION AND METHODS FOR
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MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
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GENERAL INFORMATION: Michael G.
APPLICANT: Zwick, Michael G.
Edington, Brent E.
McSwiggen, James A.
McTho, Patricia Ann Owens
Skokut, Thomas A.
Young, Scott A.
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                                                                                         APPLICANT: Hirschhorn, Joel N.
APPLICANT: Hirschhorn, Joel N.
APPLICANT: Hirschhorn, Joel N.
APPLICANT: Kaplan, Paul
APPLICANT: Kaplan, Paul
APPLICANT: Lander, Eric S.
APPLICANT: Lander, Eric S.
APPLICANT: Lockhart, David J.
TITLE OF INVENTION: UNIVERSAL ARRAYS
FILE REFRENCE: 2825.1016-007
CURRENT APPLICATION NUMBER: US 60/126,473
PRIOR APPLICATION NUMBER: US 60/126,473
PRIOR PILING DATE: 1999-03-26
PRIOR PILING DATE: 1999-06-23
PRIOR FILING DATE: 1999-06-23
PRIOR PILING DATE: 2000-03-27
NUMBER OF SEQ ID NOS: 590
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 444
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 1263
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: Template sequence US-10-730-771-444
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                           Sequence 444, Application US/10730771
Publication No. US20050074787A1
GENERAL INFORMATION:
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STATE: California
COUNTRY: U.S.A.
ZIP: 90071-2066
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US-09-877-478-1053
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                               APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Blatt, Larry
APPLICANT: Blatt, Larry
APPLICANT: Chowrita, Bharat
APPLICANT: Chowrita, Bharat
APPLICANT: Haeberli, Pete
ITITE OF INVENTION: Method and Reagent for the Inhibition of NOGO Gene
FILE REFERENCE: MBHB00, 878-A, (400/011)
CURRENT FILING DATE: 2001-02-09
FRIOR PPLICATION NUMBER: US 60/181,797
PRIOR FILING DATE: 2000-02-11
NUMBER OF SEQ ID NOS: 6679
SOCTWARE: Patentin version 3.0
SEQ ID NO S8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WESULT, 318
US-09-780-53A-59/C
; Sequence 59, Application US/09780533A
; Publication No. US2030060611A1
; GENERAL INFORMATION:
; APPLICANT: Ribozyme Pharmaceuticals, Inc.
; APPLICANT: Blatt, Larry
; APPLICANT: Chowrira, Bharat
; APPLICANT: Chowrira, Bharat
; APPLICANT: McSw4ggen, Jim
; APPLICANT: McSw4ggen, Jim
; APPLICANT: McSw4ggen, Jim
; APPLICANT: Chowrira, Bharat
; TITLE OF INVENTION: Mcthod and Reagent for the Inhibition of NOGO Gene
; TITLE OF INVENTION: Mcthod and Reagent
; CURRENT APPLICATION NUMBER: US/09/780, 533A
; CURRENT FILING DATE: 2000-02-01
; PRIOR APPLICATION NUMBER: US 60/181,797
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
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Sequence 1810, Application US/09780533A
Publication No. US20330060611A1
SEMERAL INFORNATION:
APPLICANT: Blatt, Larry
APPLICANT: McSwiggen, Jim
APPLICANT: Chowira, Bharat
APPLICANT: Haeberli, Pete
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   363 GCCGCAGGAAGAGAA 378
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Publication No. US20030060611A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 6679
SOFWARE: Patentin version 3.0
SEQ ID NO 59
EMOTH: 17
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                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Homo sapiens
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ORGANISM: Homo sapiens
US-09-780-533A-59
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Best Local Similarity
Matches 14; Conserv
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Matches 14; Conserv
                        GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                           TYPE: RNA
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APPLICANT: Ribosyme Pharmaceuticals, Inc.
APPLICANT: Ribosyme Pharmaceuticals, Inc.
APPLICANT: Blatt, Larry
APPLICANT: Blatt, Larry
APPLICANT: McSwiggen, Jim
APPLICANT: McSwiggen, Jave
TITLE OF INVENTION: Method and Reagent for Inhibiting Hepatitis B Virus Replication
FILE REFERENCE: MBHB00-845-H (400/029)
CURRENT APPLICATION NUMBER: US/09/877,478
CURRENT FILING DATE: 2001-12-31
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TITLE OF INVENTION: Method and Reagent for the Inhibition of NOGO Gene FILE REFERENCE: MBHBOO, 878-A (400/011)
CURRENT APPLICATION WUMBER: US/09/780, 533A
CURRENT FILING DATE: 2001-02-09
PRIOR APPLICATION NUMBER: US 60/181,797
PRIOR FILING DATE: 2000-02-11
NUMBER OF SEQ ID NOS: 6679
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PRIOR APPLICATION NUMBER: US 07/882,712
PRIOR PILING DATE: 1992-05-14
PRIOR PILING DATE: 1992-05-14
PRIOR FILING DATE: 2000-13-20
PRIOR FILING DATE: 2000-03-20
PRIOR PILING DATE: 2000-03-20
PRIOR APPLICATION NUMBER: US 09/636,385
PRIOR PILING DATE: 2000-08-09
PRIOR PILING DATE: 1094-02-07
PRIOR PILING DATE: 1994-02-07
PRIOR PILING DATE: 1994-02-07
PRIOR PILING DATE: 1995-05-04
PRIOR APPLICATION NUMBER: US 08/433,993
PRIOR PILING DATE: 1995-05-04
PRIOR PILING DATE: 1995-05-04
PRIOR PILING DATE: 1995-05-04
PRIOR FILING DATE: 1995-05-04
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US-09-848-754A-1036/c
; Sequence 1036, Application US/09848754A
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Sequence 1053, Application US/09877478
; Publication No. US20030068301A1
; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; ORGANISM: Homo sapiens
US-09-780-533A-1810
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Sequence 1039, Application US/09848754A
Publication No. US20030073207A1
Publication No. US20030073207A1
Publication No. US20030073207A1
Publication No. US20030073207A1

APPLICANT: Ribozyme Pharmaccuticals, Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Levels of Epidermal Growth Factor Receptors
FILE REFERENCE: MBHB00-958-I (400/018)
CURRENT APPLICATION NUMBER: US/09/948,754A
NUMBER OF SEQ ID NOS: 9645
SOFTWARE: Patentin version 3.0
SEQ ID NOS: 9645
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Publication No. US20030073207A1

GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Levels of Epidermal Growth Factor Receptors
FILE REFERENCE: MBH800-958-1 (400/018)
CURRENT APPLICATION NUMBER: US/09/848,754A

NUMBER OF SEQ ID NOS: 9645
SOFTWARE: Patentin version 3.0
SEQ ID NO 1653
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87.5%; Pred. No. 2.7e+02;
iive 0; Mismatches 2; Indels
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2.7e+02;
                 87.5%; Pred. No.
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Publication No. US20030087847A1
GENERAL INCOMPATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         199 CCCTCCCGGGGACCTG 214
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Holman, Patricia
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Matches 14; Conservative
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US-09-848-754A-1039
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US-09-848-754A-1653
              Best Local Similarity
Matches 14; Conserv
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Matches 14; Conserv
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US-09-776-474-7/c
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APPLICANT:
APPLICANT:
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                                                          APPLICANT: Ribozyme Pharmaceuticals, Inc.
TITLE OF INVENTION: Eargymatic Nucleic Acid Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Levels of Enidermal Growth Factor Receptors
PILE REFERENCE: MBHB00-958-I (400/018)
CURRENT APPLICATION WINBER: US/09/848,754A
CURRENT FILING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 9645
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Publication No. US20030073207A1
Publication No. US20030073207A1
ARDEAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate
FILE REFERENCE: MBHB00-558-I (400/018)
CURRENT APPLICATION NUMBER: US/09/848,754A
NUMBER OF SEQ ID NOS: 9645
SOFTWARE: Patentin version 3.0
SEQ ID NO 1037
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Sequence 1038, Application US/09848754A

Publication No. US20030073207A1

GENERAL INFORMATION:

APPLICANT: Ribozyme Pharmaceuticals, Inc.

TITLE OF INVENTION: Levels of Epidermal Growth Factor Receptors

TITLE OF INVENTION: Levels of Epidermal Growth Factor Receptors

FILE REFERENCE: MBHB00-958-I (400/018)

CURRENT APPLICATION NUMBER: US/09/848, 754A

CURRENT PILIOR DATE: 2001-05-03

NUMBER OF SEQ ID NOS: 9645

SOFTWARE: Patentin version 3.0
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2.8%; Score 12.8; DB 1;
Best Local Similarity 87.5%; Pred. No. 2.7e+02;
Matches 14; Conservative 0; Mismatches 2;
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                   US20030073207A1
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US-09-848-754A-1036
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ORGANISM: Homo sapiens
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ORGANISM: Homo sapiens
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US-09-848-754A-1037/c
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                   Publication No.
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LENGTH: 17
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LENGTH: 17
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US-09-930-423-335/c

1 Sequence 335, Application US/09930423

1 Sequence 335, Application US/09930423

1 Sequence 335, Application US/09930423

2 Sequence 335, Application US. US2030092003A1

3 SEPLICANT: Ribozyme Pharmaceuticals, Inc.

3 APPLICANT: Blact, Larry

4 APPLICANT: McSwigeen, Jim

5 TITLE OF INVENTION: Method and Reagent for the Treatment of Alzheimer's Disease

6 TITLE OF INVENTION NUMBER: US/09/930, 423

7 CURRENT APPLICATION NUMBER: US/09/930, 423

8 SOFTWARE: PALENTING DATE: 2001-08-15

9 SEQ ID NOS: 4553

9 SEQ ID NO 335

1 TYPE: NA

1 TYPE: NA

1 ORGANISM: Homo Sapiens

US-09-930-423-335
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Publication No. US20030092003A1
Publication No. US20030092003A1
Publication No. US20030092003A1
APPLICANT: Blact, Larry
APPLICANT: Blact, Larry
TITLE OF INVENTION: Method and Reagent for the Treatment of Alzheimer's Disease
TITLE OF INVENTION: Method and Reagent for the Treatment of Alzheimer's Disease
CURRENT APPLICATION UNBER: US/09/930,423
CURRENT FILING DATE: 2001-08-15
NUMBER OF SEQ ID NOS: 4553
SOFTWARE: Patentin version 3.0
LENGTH: 17
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2.8%; Score 12.8; DB 1;
Best Local Similarity 87.5%; Pred. No. 2.7e+02;
Matches 14; Conservative 0; Mismatches 2;
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                                                                                                          Query Match
Best Local Similarity 87.5%
Matches 14, Conservative
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                                        ; ORGANISM: Homo Sapiens
US-09-930-423-334
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; ORGANISM: Homo Sapiens
US-09-930-423-1159
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US-09-930-423-1470
                      TYPE: RNA
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APPLICANT: Fattaey, Ali
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: McSwiggen, Jim
TITLE OF INVENTION: Mcthod and Reagent for the Inhibition of Checkpoint Kinase-1 (CHK
TITLE OF INVENTION: Enzyme
TITLE OF INVENTION: Enzyme
TITLE OF INVENTION: Enzyme
TITLE OF INVENTION: Enzyme
CURRENT APPLICATION NUMBER: US 60/176,474
CURRENT FILING DATE: 2001-02-02
PRIOR FILING DATE: 2000-03-02
NUMBER OF SEQ ID NOS: 2992
SOFTWARE: PatentIn version 3.0
ENGTH: 17
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Publication No. US20030092003A1
Publication No. US20030092003A1
Publication No. US20030092003A1
APPLICANT: Blatt, Larry
APPLICANT: Blatt, Larry
APPLICANT: Matchod and Reagent for the Treatment of Alzheimer's Disease
FILE REPERENCE: MBHB00,918-A 400/027
CURRENT PRING DATE: 2001-08-15
NUMBER OF SEQ ID NOS: 4553
SOFTWARER: Patentin version 3.0
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Publication No. US20030092003A1
GENERAL INFORMATION:
APPLICANT: Bibozyme Pharmaceuticals, Inc.
APPLICANT: Blatt, Larry
APPLICANT: Machod and Reagent for the Treatment of Alzheimer's Disease
TITLE OF INVENTION: Mchod and Reagent for the Treatment of Alzheimer's Disease
CURRENT FILING DATE: 2001-08-15
NUMBER OF SEQ ID NOS: 4553
SOFTWARER Patentin version 3.0
SEQ ID NO 334
LENGTH: 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         272 CTTCTCCGGAGGCACC 287
                                                                                                                                                                                                                                                                                                                TYPE: RNA
ORGANISM: Artificial Sequence
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CORGANISM: Homo Sapiens
US-09-930-423-333
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LENGTH: 17
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Sequence 3639, Application US/09740332
Publication No. US20030125270A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals Inc.
TITLE OF INVENTION: Hepatitis C Virus Infection
TITLE OF INVENTION: Hepatitis C Virus Infection
TITLE OF INVENTION: Hepatitis C Virus Infection
CURRENT APPLICATION WUMBER: US/09/740,332
CURRENT PILING DATE: 2001-03-26
NUMBER OF SEQ ID NOS: 9704
SOFTWARE: Patentin version 3.0
SEQ ID NO 3639
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Sequence 3755, Application US/09740332

Publication No. US20030125270A1

GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Hepatitis C Virus Infection
TITLE OF INVENTION: Hepatitis C Virus Infection
TITLE OF INVENTION: HOPELIS C VIRUS INFECTION
TITLE OF INVENTION: BAPILICATION NUMBER: US/09/740,332
CURRENT FILING DATE: 2001-03-26
NUMBER OF SEQ ID NOS: 9704
SOFTWARE: PATENTIN Version 3.0

SEQ ID NO 3755

**LENGTH: 17
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                                                                                                                                                                                              ; OTHER INFORMATION: oligonucleotide substrate US-09-740-332-1425
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  SOFTWARE: Patentin version 3.0
SEQ ID NO 1425
LENGTH: 17
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ORGANISM: artificial sequence
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ORGANISM: artificial sequence
                                                                         TYPE: RNA
ORGANISM: artificial sequence
                                                                                                                                                                                                                                               Query Match
Best Local Similarity 75.0%,
-hea 12; Conservative
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                                                                                                                                                 NAME/KEY: misc_feature LOCATION:
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US-09-740-332-3639/c
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US-09-740-332-1425

US-09-740-332-1425

Sequence 1425, Application US/09740332

Publication No. US20030125270A1

GENERAL INFORMATION:

APPLICANT: RIDEZWHE Pharmaceuticals Inc.

TITLE OF INVENTION: Hepatitis C Virus Infection

FILE REFERENCE: RP1 400/003

FILE REFERENCE: RP1 400/003

CURRENT APPLICATION UMBER: US/09/740,332

CURRENT FILING DATE: 2001-03-26

NUMBER OF SEQ ID NOS: 9704
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Lawrence Blatt
APPLICANT: Lawrence McSwiggen
TITLE OF INVENTION: Wethod and Reagent for the Inhibition of NOGO and NOGO Receptor (
FILE REFERENCE: MBHB00-878-C (400/017)
CURRENT FILING DATE: 2001-04-05
PRIOR APPLICATION NUMBER: 09/780,533
PRIOR FILING DATE: 2001-02-09
PRIOR FILING DATE: 2000-02-11
NUMBER: OF SEQ ID NOS: 2617
SOFTWARE: Patentin version 3.0
SEQ ID NO 899
                                                                  APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Blatt, Larry
APPLICANT: Blatt, Larry
APPLICANT: Blatt, Larry
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Method and Reagent for the Treatment of Alzheimer's Disease
FILE REFERENCE: MBHB00,918-A 400/027
CURRENT APPLICATION NUMBER: US/09/930,423
CURRENT FILING DATE: 2001-08-15
NUMBER OF SEQ ID NOS: 4553
SOFTWARE: Patentin version 3.0
SEQ ID NO 1470
LENGTH: 17
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2.8%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 2.7e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
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Best Local Similarity 75.0%; Pred. No. 2.7e+02;
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APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: Lawrence Blatt
Sequence 1470, Application US/09930423
Publication No. US20030092003A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        263 GGCCCGGGCTTCTCC 278
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                                                                                                                                                                                                                                                                                                                                             TYPE: RNA
ORGANISM: Homo Sapiens
US-09-930-423-1470
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; ORGANISM: Homo sapiens
US-09-827-395A-899
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Sequence 334, Application US/09745237A
Publication No. US20030143708A1
GENERAL INFORMATION:
APPLICANT: Blatt, Larry
APPLICANT: Blatt, Larry
APPLICANT: Blatt, Larry
APPLICANT: MCSWiggen, Jim
TILLE OF INVENTION: McHood and Reagent for the Treatment of Alzheimer's Disease
CURRENT APPLICATION NUMBER: US/09/745,237A
CURRENT APPLICATION NUMBER: US/09/745,237A
CURRENT FILING DATE: 2002-04-15
NUMBER OF SEQ ID NOS: 4550
SOFTWARE: Patentin version 3.0
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Publication No. US20030143708A1

GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Blatt, Larry
APPLICANT: McSwiggen, Jim
FILE OF INVENTION: McHableo.918-A)
CURRENT FILMS DATE: 2002-04-15
CURRENT APPLICATION NUMBER: US/09/745,237A

NUMBER OF SEQ ID NOS: 4550
SOFTWARE: Patentin version 3.0
SOFTWARE: Patentin version 3.0
SOFTWARE: Patentin version 3.0
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US-09-745-237A-1159/C
Sequence 1159, Application US/09745237A
Sequence 1159, Application US/09745237A
Sequence 1159, Application US/09043708A1
Sequence 1159, Application Volument of Sequence 1159, Applicant in Control of Sequence 1159, Applicant in Machigan, Jim
TITLE OF INVENTION: Method and Reagent for the Treatment of Alzheimer's Disease
FILE REPERBRES. 400/07 (MBHBO0-918-A)
CURRENT APPLICATION NUMBER: US/09/745,237A
CURRENT FILING DATE: 2002-04-15
NUMBER OF SEQ ID NOS: 4550
SOFTWARE: Patentin version 3.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 2.8%; Score 12.8; DB 1; Best Local Similarity 87.5%; Pred. No. 2.7e+02; Matches 14; Conservative 0; Mismatches 2;
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US-09-745-237A-334
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ORGANISM: Homo sapiens
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Best Local Similarity
Matches 14; Conserv
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LENGTH: 17
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Publication No. US20030134806A1

GENERAL INFORMATION:

APPLICANT: Ribozyme Pharmaceuticals, Inc.

APPLICANT: Jarvis, Thale

APPLICANT: Von Carlowitz, Ira

APPLICANT: Hamblin, Paul

APPLICANT: Bilis, Jonathan

TITLE OF INVENTION: Method and Reagent for the Inhibition of Grb-2-related with Inser

TITLE OF INVENTION: (GRID) Gene

FILE REFERENCE: MBHB00-901-A (400/013)

CURRENT FILING DATE: 2001-02-23

NUMBER OF SEQ ID NOS: 2304

SEQ ID NOS: 2304

SEQ ID NO 31

LENGTH: 17
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BUBLICALION NO. 10220030143708A1

GENERAL INFORMATION:

APPLICANT: Blacty
APPLICANT: Blatt, Larry
APPLICANT: McSwiggen, Jim

TITLE OF INVENTION: Method and Reagent for the Treatment of Alzheimer's Disease
FILE REFERENCE: 400/007 (MBHB00-918-A)

CURRENT FILING DATE: 2002-04-15

NUMBER OF SEQ ID NOS: 4550
SOFTWARER: PatentIn version 3.0

SEQ ID NO 333

LENGTH: 17
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                                            Query Match

2.8%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 2.7e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
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Best Local Similarity 87.53
Matches 14; Conservative
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Matches 14; Conservative
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; ORGANISM: Homo sapiens
US-09-745-237A-333
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US-09-745-237A-333/c
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US-09-740-332-3755
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LENGTH: 17

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Sequence 3639, Application US/09817879

Publication No. US20030171311A1

GENERAL INFORMATION:

GENERAL INFORMATION:

TITLE OF INVENTION: Hepatitis C Virus Infection

TITLE OF INVENTION INVERSE: US/09/817,879

CURRENT FILING DATE: 2001-03-26

NUMBER OF SEQ ID NOS: 9703

SEQ ID NO 3639

LENGTH: 17
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US-09-817-879-3755/C
US-09-817-879-3755/C
Sequence 3755, Application US/09817879
Sequence 3755, Application US/09817879
Sequence 3755, Application US/09817811A1
GENERAL INFORMATION: Enzymatic Inc.
TITLE OF INVENTION: Hepatitis C Virus Infection
CURRENT APPLICATION UNMER: US/09/817,879
CURRENT FILING DATE: 2001-03-26
NUMBER OF SEQ ID NOS: 9703
SOFTWARE: Patentin version 3.0
SEQ ID NO 3755
LENGTH: 17
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87.5%; Pred. No. 2.7e+02;
tive 0; Mismatches 2;
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Best Local Similarity 87.5%; Pred. No. 2.7e+02;
Matches 14; Conservative 0; Mismatches 2;
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COTHER INFORMATION: Oligonucleotide substrate
US-09-817-879-3639
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; OTHER INFORMATION: oligonucleotide substrate US-09-817-879-3755
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Sequence 738, Application US/10060895A

Publication No. US20030104403A1

GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          208 GGACCTGCGGCGGGTC 223
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ORGANISM: artificial sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: RNA
ORGANISM: artificial sequence
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Best Local Similarity 87.51
Matches 14; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY: misc_feature
                                                                            US-09-817-879-3639/c
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Publication No. US20030171311A1

GENERAL INFORMATION:

APPLICANT: Ribozyme Pharmaceuticals Inc.

TITLE OF INVENTION: Hepartitis C virus Infection

TITLE OF 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Blatt, Larry
APPLICANT: McSwiggen, Jim
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: McHod and Reagent for the Treatment of Alzheimer's Disease
FILE REFERENCE: 400/007 (MBHB00-918-A)
CURRENT APPLICATION NUMBER: US/09/745,237A
CURRENT FILING DATE: 2002-04-15
NUMBER OF SEQ ID NOS: 4550
SOFTWARE: Patentin version 3.0
SEQ ID NO 1470
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2.8%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 2.7e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
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87.5%; Pred. No. 2.7e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 1470, Application US/09745237A Publication No. US20030143708A1 GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                242 CCCCGCCTGGAGGCCG 257
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                                                                                                                                                                                                                                                                                 12 GGTGGCCTGGGAGGG 27
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Matches 14; Conservative
TYPE: RNA
CORGANISM: Homo sapiens
US-09-745-237A-1159
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: RNA
GRGANISM: Homo sapiens
US-09-745-237A-1470
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US-09-745-237A-1470
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US-09-817-879-1425
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APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: NGSWiggen, Jim
TITLE OF INVENTION: Nucleic acid treatment of diseases or conditions related to level:
TITLE OF INVENTION: HEBZ.
FILE REPERENCE: MBHB01-1653-A (400/014)
CURRENT APPLICATION NUMBER: US/10/163,552
CURRENT FILING DATE: 2002-06-06
NUMBER OF SEQ ID NOS: 1997
SSPTWARE: Patentin version 3.0
SSPTWARE: Patentin version 3.0
LENGTH: 17
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Publication No. US20030105051A1
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Roswiggen, Jim
TITLE OF INVENTION: Nucleic acid treatment of diseases or conditions related to level
TITLE OF INVENTION: HERS
FILE REFERENCE: MBHB01-1653-A (400/014)
CURRENT APPLICATION NUMBER: US/10/163,552
CURRENT FILING DATE: 2002-06-06
NUMBER OF SEQ ID NOS: 1997
SOFTWARE: PatentIn version 3.0
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2.8%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 2.7e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
                       PRIOR FILING DATE: 2001-08-30
NUMBER OF SEQ ID NOS: 1682
SOFTWARE: Acomica Sequence Listing Engine
SEQ ID NO 739
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; Sequence 20, Application US/10163552; Publication No. US20030105051A1; GENERAL INFORMATION:
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                                                                                                                                                                                                        TYPE: DNA
CORGANISM: Homo sapiens
US-10-060-895A-739
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; TYPE: RNA
; ORGANISM: Homo sapiens
US-10-163-552-20
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ORGANISM: Homo sapiens
US-10-163-552-869
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Best Local Similarity
Matches 14; Conserv
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US-10-163-552-869/c
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LENGTH: 17
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Sequence 739, Application US/10060895A;
Publication No. US20030104403A1
GENERAL INFORMATION:
APPLICANT: Chang, Jian
APPLICANT: Chang, Jian
APPLICANT: Chung-Tuong
TITLE OF INVERTION: Chung-Tuong
TITLE OF INVERTION: UNIVERS: US/10/060,895A
CURRENT FILING DATE: 2001-01-30
FRICE REPERING: PED156
FRICE REPERING: POOL-06-10
FRICE REPLICATION NUMBER: PCT/US01/0066
FRICE REPLICATION NUMBER: PCT/US01/0066
FRICE PRING DATE: 2001-01-30
FRICE RILING DATE: 2001-01-30
FRICE RELING DATE: 2001-01-30
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2.8%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 2.7e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
                                                                                                                                              CURRENT FILING DATE: 2002-06-10

PRIOR APPLICATION NUMBER: PCT/USO1/00666

PRIOR PILING DATE: 2001-01-30

PRIOR APPLICATION NUMBER: PCT/USO1/00665

PRIOR PILING DATE: 2001-01-30

PRIOR PILING DATE: 2001-01-30
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ORGANISM: Homo sapiens
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APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: McSwiggen, James
TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or Conditions Related to Level
FILE REFERENCE: 400/057 (MBHB01-1158-A)
CURRENT APPLICATION NUMBER: US/10/238,700
CURRENT FILING DATE: 2002-09-18
PRIOR FILING DATE: 2002-05-29
PRIOR FILING DATE: 2001-09-10
NUMBER OF SEQ ID NOS: 4666
SOFTWARE: Patentin version 3.0
SEQ ID NO 2821
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APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: McSwiggen, James
TITLE OF INVENTION: UNICLE: A Acid Treatment of Diseases or Conditions Related to Level
FILE REFERENCE: 400/057 (MBHB01-1158-A)
CURRENT APPLICATION NUMBER: US/10/238,700
CURRENT FILING DATE: 2002-09-18
PRIOR FILING DATE: 2002-05-29
PRIOR PILING DATE: 1002-05-29
PRIOR PILING DATE: 2001-09-10
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                                                                                                                                                                                                                                                                                                                               2; Indels
                                                                                                                                                                                                                                                                           2.8%; Score 12.8; DB 1;
87.5%; Pred. No. 2.7e+02;
trive 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 12.8; DB 1;
Pred. No. 2.7e+02;
2; Mismatches 2;
                            60/318,471
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Sequence 2821, Application US/10238700
; Publication No. US20030153521A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; Sequence 3508, Application US/10238700 ; Publication No. US20030153521A1 ; GENERAL INFORMATION:
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PRIOR FILING DATE: 2002-05-29
PRIOR APPLICATION NUMBER: US 60
PRIOR FILING DATE: 2001-09-10
NUMBER OF SEQ ID NOS: 4666
SOFTWARE: PATENTIN VERSION 3.0
SEQ ID NO 2700
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SOFTWARE: PatentIn version 3.0
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Best Local Similarity 75.0%;
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Best Local Similarity 87.5°
Matches 14, Conservative
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; ORGANISM: Homo sapiens
US-10-238-700-2700
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; ORGANISM: Homo sapiens
US-10-238-700-2821
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ORGANISM: Homo sapiens
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LENGTH: 17
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                                                            RESULT 349
US-10-156-306-4930
Sequence 4930, Application US/10156306
Sequence 4930, Application US/10156306
Sequence 4930, Application No. US20030119017A1
Publication No. US20030119017A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: MESWYIGHTON:
ITILE OF INVENTION: Levels of IKK-Gamma and PKR
FILE REPERENCE: MHB01-664-A (400/050)
CURRENT PRILING DATE: 2002-05-28
NUMBER OF SEQ ID NOS: 8013
SOFTWARE: Patentin version 3.0
SEQ ID NO 4930

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***CHARMATE: Patentin version 3.0
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; Publication No. US20030119017A1
; GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: McSwiggen, James
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Levels of IKK-Gamma and PKR
FILE REFERENCE: MBHB01-664-A (400/050)
CURRENT APPLICATION NUMBER: US/10/156,306
CURRENT APPLICATION NUMBER: US/10/156,306
CURRENT FILING DATE: 2002-05-28
SOUTHWER OF SEQ ID NOS: 8013
SOFTWARE: Patentin version 3.0
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Publication No. US20030153521A1
APDEDICANT: NFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: McSwiggen, James
TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or Conditions Related to Level
CURRENT APPLICATION NUMBER: US/10/238,700
CURRENT FILING DATE: 2002-09-18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      249 TGGAGGCCGCGGTCGG 264
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              246 GCCTGGAGGCCGCGGT 261
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  17 GAGGAGGGTGGGCTG 2
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Matches 12; Conservative
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Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; TYPE: RNA
; ORGANISM: Homo sapiens
US-10-156-306-4930
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: RNA
ORGANISM: Homo sapiens
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US-10-238-700-2700/c
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US-10-156-306-6942
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Sequence 1249, Application US/10230006

Subblication No. US20030191077A1

GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Resnaugh, Kathy
APPLICANT: Resnaugh, Kathy
TITLE OF INVENTION: METHOD AND REAGENT FOR THE TREATMENT OF ASTHWA AND ALLERGIC CONDITION ENTRY APPLICATION NUMBER: US/10/230,006
CURRENT APPLICATION NUMBER: US/10/230,006
CURRENT FILING DATE: 2002-11-18
PRIOR APPLICATION NUMBER: US 60/315,315
PRIOR APPLICATION NUMBER: US 60/315,315
PRIOR PILING DATE: 2001-08-28
NUMBER OF SEQ ID NOS: 2678
SOFTWARE PARENT APPLICATION NUMBER OF SEQ ID NOS: 2679
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Sequence 1284, Application US/10230006

Sequence 1284, Application US/10230006

Sequence 1284, Application US/10230006

GENERAL INFORMATION:

APPLICANT: Ribozyme Pharmaceuticals, Inc.

APPLICANT: Rosnaugh, Xathy

APPLICANT: McSwiggen, Jim

TITLE OF INVENTION: METHOD AND REAGENT FOR THE TREATMENT OF ASTHMA AND ALLERGIC CONDI

FILE REFERENCE: 400/056 (MBHB01-1110)
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   PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/USO1/00669
PRIOR FILING DATE: 2001-01-30
PRIOR PLING DATE: 2001-01-30
PRIOR PLING DATE: 2001-01-30
PRIOR PLING DATE: 2001-01-30
PRIOR FILING DATE: 2001-05-23
PRIOR FILING DATE: 2001-05-23
PRIOR FILING DATE: 2001-01-010
NUMBER OF SEQ ID NOS: 4162
SOFTWARE: Acomica Sequence Listing Engine
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US-10-061-201-716
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US-10-230-006-1249
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
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                                                        2.8%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 2.7e+02; tive 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION:
TITLE OF INVENTION: HUMAN POSH-LIKE PROTEIN 1
FILE REFERENCE: PB0178
CURRENT APPLICATION NUMBER: US/10/061,201
CURRENT PILING DATE: 2002-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR PILING DATE: 2001-10-10
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US-10-061-201-716

Sequence 716, Application US/10061201

Publication No. US20030166229A1

GENERAL INFORMATION:

APPLICANT: Shannon, Mark

TITLE OF INVENTION: HUMAN POSH-LIKE PROTEIN 1

FILE REFERENCE: PB0178

CURRENT APPLICATION NUMBER: US/10/061,201

CURRENT FILING DATE: 2002-01-30

PRIOR APPLICATION NUMBER: PCT/US01/00666

PRIOR APPLICATION NUMBER: PCT/US01/00667

PRIOR PILING DATE: 2001-01-30

PRIOR PILING DATE: 2001-01-30

PRIOR PILING DATE: 2001-01-30

PRIOR APPLICATION NUMBER: PCT/US01/00667
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SEQ ID NO 715
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Publication No. US20030166229A1
GENERAL INFORMATION:
                                                                                                                                                                           226 CTGCCCAGCCCCGAA 241
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                                                 Query Match
Best Local Similarity 87.5:
Matches 14; Conservative
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US-10-238-700-3508
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Sequence 751, Application US/10307005

Sequence 751, Application US/10307005

Publication No. US2003023620841

GENERAL INDOFMATION:

APPLICANT: University of Delaware

APPLICANT: Eric B. Kmiec

APPLICANT: Michael C. Rice

APPLICANT: Michael C. Rice

APPLICANT: Michael C. Rice

APPLICANT: Wichael C. Rice

APPLICANT: Wichael C. Rice

APPLICANT: Unigsup Kim

TITLE OF INVENTION: Using Modified Single Stranded Oligonucleotides

TITLE OF INVENTION: Using Modified Single Stranded Oligonucleotides

TITLE OF INVENTION: Using Modified Single Stranded Oligonucleotides

FILE REPERBENCE: Napro/009 PCT

CURRENT APPLICATION NUMBER: US/10/307,005

PRIOR PILING DATE: 2000-06-01

PRIOR PILING DATE: 2000-06-01

PRIOR PILING DATE: 2000-06-01

PRIOR PILING DATE: 2000-10-30

PRIOR PILING DATE: 2000-10-30

PRIOR FILING DATE: 2000-10-30

PRIOR PILING DATE: 2000-10-30

PRIOR FILING DATE: 2000-10-30

PRIOR FILING DATE: 2000-10-30

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                                                                                                                                                                                                                                                                                                                                                                                                                                     ; OTHER INFORMATION: Description of Artificial Sequence:capture US-10-297-068-1071
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          TITLE OF INVENTION: KIT AND METHOD FOR DETERMINING HLA TYPES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
2.8%; Score 12.8; DB 1; Length 1
Best Local Similarity 87.5%; Pred. No. 2.7e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
                              FILE REFERENCE: 1314OP1174
CURRENT APPLICATION NUMBER: US/10/297,068
CURRENT FILING DATE: 2002-11-27
FRIOR APPLICATION NUMBER: UP 2000-164798
PRIOR FILING DATE: 2000-06-01
NUMBER OF SEQ ID NOS: 1298
SOFTWARE: PatentIn Ver. 2.1
LENGTH: 17
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APPLICANT: University of Delaware APPLICANT: Eric B. Kmiec APPLICANT: Howard B. Gamper
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           410 CTGAGCTGTGGGACGT 425
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                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
ORGANISM: Artificial Seguence
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Matches 14; Conserv
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US-10-307-005-752/c
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Sequence 899, Application US/10430882

Publication No. US20030203870A1

GENERAL INFORMATION:

APPLICANT: Lawrence Blatt

APPLICANT: Lawrence Blatt

APPLICANT: Bharat ChowTira

APPLICANT: All ChowTira

FRIOR APPLICATION NUMBER: 09/780,533

PRIOR FILING DATE: 2001-02-09

PRIOR FILING DATE: 2001-02-09

PRIOR FILING DATE: 2001-02-09

PRIOR FILING DATE: 2000-02-11

PRIOR APPLICATION NUMBER: PCT/US02/10512

SEQUENCE: Bacentin version 3.0

SEQUENCE: Bacentin version 3.0
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                                                                                                                                                                                                                                                                                                                                                                                               Query Match
2.8%; Score 12.8; DB 1; Length 17;
Best Local Similarity 68.8%; Pred. No. 2.7e+02;
Matches 11; Conservative 3; Mismatches 2; Indels
CURRENT APPLICATION NUMBER: US/10/230,006
CURRENT FILING DATE: 2002-11-18
PRIOR APPLICATION NUMBER: US 60/315,315
PRIOR FILING DATE: 2001-08-28
NUMBER OF SEQ ID NOS: 2678
SOFTWARE: Patentin version 3.0
SEQ ID NO 1284
LENGTH: 17
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Publication No. US20030228585A1
GENERAL INFORMATION:
APPLICANT: INOKO, Hidetoshi
APPLICANT: KAGIYA, Taeko
APPLICANT: ICHIHARA, TATSUO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  KAGIYA, Taeko
ICHIHARA, Tatsuo
Matsumura, Yoshiyuki
MORIYA, Shogo
NISHIDA, Michio
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        182 GCTGCTGCCCGTTCG 197
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                                                                                                                                                                                                                                                                    TYPE: RNA
GORGANISM: Homo sapiens
US-10-230-006-1284
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; ORGANISM: Homo sapiens
US-10-430-882-899
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APPLICANT:
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USACHELLANDER OF STATA (C 18204017) 18674

1 Sequence 8374, Application US/10138674

2 Fublication No. US20040077565A1

3 Fublication No. US20040077565A1

3 Fublicant: Ribozyme Pharmaceuticals, Inc.

4 APPLICANT: Ribozyme Pharmaceuticals, Inc.

5 APPLICANT: Bacobed, Jam

5 APPLICANT: Stinchcomb, Dan

6 APPLICANT: Stinchcomb, Dan

7 APPLICANT: Bacobed, Jaime

7 TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor

7 TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor

8 FILE REFERENCE: MBHB00-876-N (400/049)

7 CURRENT APPLICATION NUMBER: US/10/138,674

7 CURRENT PLING DATE: 2002-05-03

8 SOFTWARE: Patentin version 3.0

7 END ID NO 8374

7 END ID NO 8374
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APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Bavco, Pam
APPLICANT: Bavco, Pam
APPLICANT: StinchComb, Jam
APPLICANT: StinchComb, Jam
APPLICANT: Escobedo, Jaime
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Merbod and Reagent for the Treatment of Diseases or Conditions Re
TITLE OF INVENTION: Merbod and Reagent for the Treatment of Diseases or Conditions Re
FILE REFERENCE: MBHB00-876-N (400/049)
CURRENT APPLICATION NUMBER: US/10/287,949A
CURRENT FILING DATE: 2003-04-11
NUMBER OF SEQ ID NOS: 20822
SEQ ID NOS: 20822
SEQ ID NOS: 20822
LENGTH: 17
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2.8%; Score 12.8; DB 1;
Best Local Similarity 87.5%; Pred. No. 2.7e+02;
Matches 14; Conservative 0; Mismatches 2;
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. Sequence 8374, Application US/10287949A
. Publication No. US20040102389A1
. GENERAL INFORMATION:
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                             363 GCCGCAGGAAGAGAA 378
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; TYPE: RNA
; ORGANISM: Homo sapiens
US-10-138-674-8374
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ORGANISM: Homo sapiens
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APPLICANT: Blatt, Larry
APPLICANT: Blatt, Larry
APPLICANT: Blatt, Larry
APPLICANT: Blatt, Larry
APPLICANT: Maxwiggen, Jin
APPLICANT: Morrisesy, Dave
FILE REFRERNCE: 400/075 (MAHBOO-484-1)
CURRENT FILING DATE: 2003-01-15
CURRENT FILING DATE: 2003-01-15
PRIOR FILING DATE: 2000-08-09
PRIOR PELING DATE: 2000-08-09
PRIOR FILING DATE: 2000-08-09
PRIOR FILING DATE: 2000-08-09
PRIOR PELING DATE: 2000-08-09
PRIOR PELING DATE: 2000-08-09
PRIOR PELING DATE: 2000-10-24
PRIOR PELING DATE: 1994-02-07
PRIOR FILING DATE: 1994-02-07
PRIOR FILING DATE: 1994-02-07
PRIOR FILING DATE: 1992-05-14
PRIOR FILING DATE: 1992-05-14
PRIOR FILING DATE: 1992-05-14
PRIOR FILING DATE: 1992-01-08
PRIOR FILING DATE: 1992-01-08
PRIOR FILING DATE: 1992-05-14
PRIOR FILING DATE: 1992-05-14
PRIOR FILING DATE: 1992-05-14
PRIOR FILING DATE: 1992-01-08
PRIOR FILING DATE: 1993-01-08
APPLICANT: Michael C. Rice
APPLICANT: Unugsup Kim
ITILE OF INVENTION: Targeted Chromosomal Genomic Alterations in Plants
ITILE OF INVENTION: Targeted Chromosomal Genomic Alterations in Plants
ITILE OF INVENTION: Using Modified Single Stranded Oligonucleotides
FILE REFERENCE: Napro/009 PCT of Single Stranded Oligonucleotides
CURRENT APPLICATION NUMBER: US/01/0307,005
PRIOR PLILNG DATE: 2000-06-01
PRIOR PLILNG DATE: 2000-06-01
PRIOR PLILNG DATE: 2000-06-01
PRIOR PLILNG DATE: 2000-06-01
PRIOR APPLICATION NUMBER: US 60/244,989
PRIOR PLILNG DATE: 2000-10-30
PRIOR PLILNG DATE: 2000-10-30
PRIOR PLILNG DATE: 2000-03-27
NUMBER OF SEQ ID NOS: 2717
SEQ ID NOS: 2717
SEQ ID NOS: 2717
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2.8%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 2.7e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
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Publication No. US20040054156A1
GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Draper, Kenneth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            410 CTGAGCTGTGGGACGT 425
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ORGANISM: Hepatitis B virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; ORGANISM: Zea mays
US-10-307-005-752
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APPLICANT: Partico, Lee
APPLICANT: Partico, Lee
APPLICANT: Kenneth, Draper
APPLICANT: Kenneth, Draper
APPLICANT: Kenneth, Draper
APPLICANT: Renneth, Draper
APPLICANT: Renneth, Draper
APPLICANT: Elisabeth, Roberta
APPLICANT: Elisabeth, Roberta
TITLE OF INVENTION: OLIGOMUCLECTIDE MEDIATED INHIBITION OF HEPATITIS B VIRUS AND HEPJ
TITLE OF INVENTION: VIRUS REPLICATION
FILLS OF INVENTION: VIRUS REPLICATION
FILLS OF INVENTION: OLIGOMOCLECTION
FILLS OF INVENTION NUMBER: US (120,203-03-04)
FRIOR FILLNG DATE: 2001-06-08
FRIOR PELICATION NUMBER: US (60/356,876
FRIOR PELICATION NUMBER: US (60/356,876
FRIOR APPLICATION NUMBER: US (60/356,580
FRIOR PELICATION NUMBER: US (60/363,124
FRIOR PELICATION NUMBER: US (60/363,124
FRIOR FILING DATE: 2000-20-20
FRIOR FILING DATE: 2000-20-10
FRIOR FILING DATE: 2000-00-10
FRIOR FILING DATE: 2000-00-1
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APPLICANT: Lawrence, Blatt
APPLICANT: Dennis, Macejak
APPLICANT: Dennis, Macejak
APPLICANT: Dennis, Macejak
APPLICANT: Dennis, Macejak
APPLICANT: Dennis, Marrissey
APPLICANT: Pamela, Pavco
APPLICANT: Partice, Lee
APPLICANT: Renneth, Draper
APPLICANT: Elisabeth, Roberts
TITLE OF INVENTION: UIRUS REPLICATION
FILE REFERENCE: 400/042US (MBHB02-249-E)
CURRENT APPLICATION NUMBER: US/10/669,841
CURRENT FILING DATE: 2003-09-23
PRIOR PELICATION NUMBER: DCT/USO2/09187
PRIOR PELICATION NUMBER: US/20/64,876
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
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Publication No. US20040127446A1
GENERAL INFORMATION:
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                                                                                                                                                             Pamela, Pavco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
Matches 14; Conserv
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                                             APPLICANT: CHOWALALL, CHOWALALL, CHOWALALL, CHOWALALL, CHOWALDEANT: APPLICANT: BLICKOMD, Dan APPLICANT: StinchComb, Dan ITILE OF INVENTION: Method and Reagent for the Inhibition of Telomerase Enzyme FILE REPERENCE: MBHB00-882-C (400/019)
CURRENT APPLICATION NUMBER: US/09/653,225
FRIOR APPLICATION NUMBER: US/09/653,225
FRIOR FILING DATE: 2000-08-31
FRIOR FILING DATE: 2000-08-31
FRIOR FILING DATE: 2000-08-31
FRIOR FILING DATE: 2000-08-31
FRIOR FILING DATE: 1999-08-31
NUMBER OF SEQ ID NOS: 5586
SSOFIWARE: PARENTIN VERBION 3.0
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SSOFIWARE: PARENTIN VERBION 3.0
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Sequence 1149, Application US/10712672

Publication No. US20040102413A1

Sequence 1149, Application US/10712672

Publication No. US20040102413A1

Sequence 1149, Application No. US20040102413A1

APPLICANT: Ribozyme Pharmaceuticals, Inc.

APPLICANT: Ribozyme Pharmaceuticals, Inc.

APPLICANT: Chowitra, Bharat

APPLICANT: Chowitra, Bharat

APPLICANT: Chowitra, Bharat

APPLICANT: McSwiggen, Jan

TITLE OF INVENTION: Method and Reagent for the Inhibition of Telomerase Enzyme

FILER REPERENCE: MBHB00-892-C (400/019)

CURRENT FILING DATE: 2003-11-13

PRIOR PELING DATE: 2000-08-31

PRIOR PPLICATION NUMBER: 60/197,769

PRIOR APPLICATION NUMBER: 60/197,769

PRIOR APPLICATION NUMBER: 60/150,713

PRIOR APPLICATION UNMBER: 60/150,713

PRIOR APPLICATION UNMBER: 60/150,713

PRIOR APPLICATION NUMBER: 60/150,713
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2.8%; Score 12.8; DB 1;
Best Local Similarity 87.5%; Pred, No. 2.7e+02;
Matches 14; Conservative 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
2.8%; Score 12.8; DB 1;
Best Local Similarity 87.5%; Pred. No. 2.7e+02;
Matches 14; Conservative 0; Mismatches 2;
APPLICANT: Ribozyme Pharmaceuticals, Inc.
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US-10-712-672-757
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APPLICANT: REINEABEH, ReDerts

TITLE OF INVENTION: OLIGONUCLECTIDE MEDIATED INHIBITION OF HEPATITIS B VIRUS AND HEPA,
TITLE OF INVENTION: VIRUS REPLICATION
TITLE OF INVENTION: VIRUS REPLICATION
FILE SEPERENCE: 400/042US (MBHBO2-249-E)
CURRENT PAPLICATION NUMBER: US/10/669,841
CURRENT PAPLICATION NUMBER: PCT/US02/09187
PRIOR APPLICATION NUMBER: PCT/US02/09187
PRIOR PLING DATE: 2001-06-08
PRIOR APPLICATION NUMBER: US 60/235,059
PRIOR APPLICATION NUMBER: US 60/337,055
PRIOR APPLICATION NUMBER: US 60/337,055
PRIOR PLING DATE: 2001-10-24
PRIOR PLING DATE: 2001-12-05
PRIOR PLING DATE: 2001-12-06
PRIOR PLING DATE: 2001-12-06
PRIOR PLING DATE: 2001-12-06
PRIOR PLING DATE: 2001-12-06
PRIOR PLING DATE: 2001-12-07
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                       PRIOR FILING DATE: 2001-03-26
PRIOR APPLICATION NUMBER: US 09/740,332
PRIOR PILING DATE: 2000-12-18
PRIOR FILING DATE: 2000-07-07
PRIOR FILING DATE: 2000-07-07
PRIOR FILING DATE: 2000-02-15
PRIOR FILING DATE: 2000-02-15
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 16207
SOFTWARE: PatentIn version 3.0
SEQ ID NO 6232
LENGTH: 17
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; Sequence 6348, Application US/10669841
; Publication No. US20040127446A1
; GENERAL INFORMATION:
; APPLICANT: Sirna Therapeutics, Inc.
; APPLICANT: Lawrence, Blatt
; APPLICANT: Dennis, Macejak
; APPLICANT: Dennis, McSwiggen
; APPLICANT: Dennis, McSwiggen
; APPLICANT: Denid, Morrissey
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ORGANISM: Artificial Sequence
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APPLICANT: Patrice, Lee
APPLICANT: Kenneth, Draper
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NAME/KEY: misc_feature
LOCATION:
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APPLICANT: Blisabeth, Diaper
APPLICANT: Elisabeth, Diaper
APPLICANT: Elisabeth, Reberts
ITILE OF INVENTION: OLICONUCLECTIDE MEDIATED INHIBITION OF HEPATITIS B VIRUS AND HEPA
ITILE OF INVENTION: VIRUS REPLICATION
FILE REPRENCE: 400/04202 (MAHBAR2-249-E)
CURRENT APPLICATION NUMBER: US/10/669,841
PRIOR APPLICATION NUMBER: US 60/296,876
PRIOR APPLICATION NUMBER: US 60/296,876
PRIOR APPLICATION NUMBER: US 60/335,059
PRIOR FILING DATE: 2001-10-24
PRIOR PLING DATE: 2001-10-24
PRIOR PLING DATE: 2001-12-05
PRIOR APPLICATION NUMBER: US 60/337,055
PRIOR APPLICATION NUMBER: US 60/337,055
PRIOR PLING DATE: 2002-12-05
PRIOR PLING DATE: 2002-12-05
PRIOR PPLICATION NUMBER: US 60/363,124
PRIOR PLING DATE: 2002-03-11
PRIOR PLING DATE: 2002-03-11
PRIOR PLING DATE: 2002-03-11
PRIOR PLING DATE: 2002-03-11
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PRIOR FILING DATE: 2001-06-08

PRIOR APPLICATION NUMBER: US 60/335,059

PRIOR APPLICATION NUMBER: US 60/337,055

PRIOR PILING DATE: 2001-10-26

PRIOR FILING DATE: 2001-10-05

PRIOR FILING DATE: 2002-02-0

PRIOR PILING DATE: 2002-03-11

PRIOR APPLICATION NUMBER: US 60/363,124

PRIOR APPLICATION NUMBER: US 60/363,124

PRIOR PILING DATE: 2001-03-61

PRIOR APPLICATION NUMBER: US 09/917,879

PRIOR PILING DATE: 2000-01-03-61

PRIOR APPLICATION NUMBER: US 09/740,332

PRIOR APPLICATION NUMBER: US 09/5011,931

PRIOR PILING DATE: 2000-02-07

PRIOR PILING DATE: 2000-02-07

PRIOR PILING DATE: 2000-02-15

Remaining Prior Application data removed - See File Wrapper or PALM.

NUMBER OF SEQ ID NOS: 16207
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2.8%; Score 12.8; DB 1; Length 17;
Best Local Similarity 75.0%; Pred. No. 2.7e+02;
Matches 12; Conservative 2; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ) OTHER INFORMATION: oligonucleotide substrate US-10-669-841-4018
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Publication No. US20040127446A1
GENERAL INFORMATION:
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APPLICANT: Lawrence, Blatt
APPLICANT: Dennis, Macejak
APPLICANT: James, McKwiggen
APPLICANT: David, Morrissey
APPLICANT: Pamela, Pavco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: RNA
ORGANISM: Artificial Sequence
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NAME/KEY: misc_feature
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LENGTH: 17
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APPLICANT: MCSAIGGEN, James
TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or Conditions Related to Level
TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or Conditions Related to Level
TITLE OF INVENTION: Ras, HERS, and HIV
FILE REFERENCE: 400/046-US (MBHB02-326-A)
CURRENT APPLICATION NUMBER: US/10/724,270
CURRENT FILING DATE: 2001-126
PRIOR APPLICATION NUMBER: US 60/184,471
PRIOR APPLICATION NUMBER: US 60/294,140
PRIOR FILING DATE: 2001-06-06
PRIOR FILING DATE: 2001-06-10
PRIOR FILING DATE: 2001-06-10
PRIOR FILING DATE: 2001-06-10
PRIOR FILING DATE: 2002-09-10
PRIOR FILING DATE: 2002-09-10
PRIOR FILING DATE: 2002-09-10
PRIOR FILING DATE: 2002-06-10
PRIOR FILING DATE: 2002-06-06
PRIOR FILING DATE: 2002-06-06
PRIOR FILING DATE: 2002-06-03
PRIOR FILING DATE: 2002-06-03
PRIOR FILING DATE: 2002-06-03
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PRIOR PRIOR PRIOR DATE: 2003-06-06-06
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PRIOR PRIOR DATE: 2003-06-06-06
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APPLICANT: McSwiggen, James
TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or Conditions Related to Level)
TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or Conditions Related to Level)
TITLE OF INVENTION: RAS, HERZ and HIV
FILE REFERENCE: 400/046-US (MBHB02-326-A)
CURRENT APPLICATION NUMBER: US/10/724,270
CURRENT FILING DATE: 2003-11-26
PRIOR APPLICATION NUMBER: US 60/318,471
PRIOR APPLICATION NUMBER: US 60/296,249
PRIOR PILING DATE: 2001-09-10
PRIOR PLING DATE: 2001-06-06
PRIOR APPLICATION NUMBER: US 60/294,140
PRIOR PILING DATE: 2001-05-06
PRIOR APPLICATION NUMBER: US 10/1338,700
PRIOR APPLICATION NUMBER: US 10/163,552
PRIOR PILING DATE: 2002-06-06
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; Sequence 1500, Application US/10724270
; Publication No. US20050080031A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         254 GCCGCGGTCGGCCCGG 269
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PatentIn version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       h 2.8%;
Similarity 87.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16 eccececión 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity v...
Best Local Similarity by...
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; TYPE: RNA
; ORGANISM: Homo sapiens
US-10-724-270-1379
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APPLICANT: Sandberg, Jennifer
APPLICANT: Sandberg, Jennifer
APPLICANT: Gordon, Gilad
APPLICANT: Gordon, Gilad
APPLICANT: Gordon, Gilad
APPLICANT: Gordon, Gilad
APPLICANT: Stinchcomb, Dan
TITLE OF INVENTION: NUCLEIC ACID BASED MODULATION OF VASCULAR ENDOTHELIAL GROWTH FACT
TITLE OF INVENTION: NUCLEIC ACID BASED MODULATION OF VASCULAR ENDOTHELIAL GROWTH FACT
TITLE OF INVENTION: NUCLEIC ACID BASED MODULATION OF VASCULAR ENDOTHELIAL GROWTH FACT
TITLE OF INVENTION: NUMBER: US 60/065,974
FRIOR PELING DATE: 1995-10-11-3
FRIOR PELING DATE: 1995-01-08
FRIOR PELING DATE: 1995-01-08
FRIOR PELING DATE: 1995-08-10
FRIOR PELING DATE: 2000-11-07
FRIOR PELING DATE: 2001-11-07
FRIOR PELING DATE: 2001-11-30
FRIOR PELING DATE: 2001-05-39
FRIOR PELING DATE: 2001-11-30
FRIOR PELING D
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                                                                                                                                                                                                               OTHER INFORMATION: Description of Artificial Sequence: Nucleic Acid
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2.8%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 2.7e+02;
Matches 14; Conservative 0; Mismatches 2: Indels
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                                                                                                                                                                                                                                                                                            NAME/KEY: misc_feature
LOCATION:
OTHER INFORMATION: oligonucleotide substrate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 3415, Application US/10712633
Publication No. US20040220128A1
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
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                                                                                     TYPE: RNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17 GGACTGGGCCCACACA 2
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ORGANISM: Homo Sapiens
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Best Local Similarity
Matches 14; Conserva
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US-10-712-633-3415/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 372
US-10-724-270-1379/c
                                                                                                                                                                                                                                                                                                                                                                                                                                    US-10-669-841-6348
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SEQ ID NO 6348
LENGTH: 17
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1.5 Sequence 4675, Application US/10724270
5. Sequence 4675, Application US/10724270
5. Sequence 4675, Application US/10724270
5. Publicarion No. US20050080031A1
6. RAPLICANT: Sirna Therapeutics, Inc.
6. APPLICANT: Mosbiggen, James
7. TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or Conditions Related to Levelt
7. TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or Conditions Related to Levelt
7. TITLE OF INVENTION: Number: 2003-11-26
7. TITLE OF INVENTION: NUMBER: 92 (Maha02-326-A)
7. CURRENT APPLICATION NUMBER: 92 (Maha02-326-A)
7. CURRENT APPLICATION NUMBER: 95 (Maha02-326-A)
7. CURRENT APPLICATION NUMBER: 95 (Maha02-326-A)
7. FRIOR APPLICATION NUMBER: 91 (Maha02-326-A)
7. FRIOR PLING DATE: 2002-06-10
7. FRIOR PLING DATE: 2002-06-10
7. FRIOR PLING DATE: 2002-06-10
7. FRIOR PLING DATE: 2003-06-31
7. FRIOR DATE: 2003-06-31
7. FRIOR DATE: 2003-06-31
7. FRIOR DATE: 2003-06-31
7. FRIOR DATE: 2003-06-31
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Sequence 5524, Application US/10724270

Publication No. US20050080031A1

GENERAL INFORMATION:

APPLICANT: Sirna Therapeutics, Inc.

APPLICANT: McSwiggen, James

TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or Conditions Related to Levels

TITLE OF INVENTION: RAS, HERZ and HIV

TITLE OF INVENTION: RAS, HERZ and HIV

FILE REFERENCE: 400/046-US (MBHB02-336-A)

CURRENT FILING DATE: 2003-11-26

FRIOR APPLICATION NUMBER: US/10/724,270

FRIOR APPLICATION NUMBER: US 60/296,249

FRIOR APPLICATION NUMBER: US 60/296,249

FRIOR APPLICATION NUMBER: US 60/296,140

FRIOR PILING DATE: 2001-05-29

FRIOR APPLICATION NUMBER: US 60/294,140

FRIOR PILING DATE: 2001-05-29

FRIOR PILING DATE: 2001-05-29

FRIOR PILING DATE: 2001-06-06

FRIOR FILING DATE: 2002-09-10

FRIOR FILING DATE: 2002-09-10
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Best Local Similarity 87.5%; Pred. No. 2.7e+02;
Matches 14; Conservative 0; Mismatches 2;
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-10-724-270-5524/c
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US-10-724-270-2187/c
| Sequence 2187, Application US/10724270
| Sequence 2187, Application US/10724270
| Publication No. US20050080031A1
| GENERAL INFORMATION: USCASOSOBORIAL
| APPLICANT: SITUATION: USCASOSOBORIAL
| TITLE OF INVENTION: NACA: Application Value
| TITLE OF INVENTION: RAS, HERZ and HIV.
| TITLE OF INVENTION: NACA: GENERAL ADDITION OF SET OF S
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PRIOR APPLICATION NUMBER: US 10/444,853
PRIOR FILING DATE: 2003-05-23
PRIOR FILING DATE: 2003-04-16
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 6810
SOFTWARE: Patentin version 3.0
SEQ ID NO 1500
LENGTH: 17
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US-10-724-270-1500
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US-10-724-270-2187
                                                                                                                                                                                                                                                                                                                                                                                         TYPE: RNA
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PRIOR APPLICATION NUMBER: US 10/157,580

PRIOR FILING DATE: 2002-05-29

PRIOR PLILING DATE: 2002-05-29

PRIOR PLILING DATE: 2002-10-23

PRIOR PLILING DATE: 2003-05-23

PRIOR PLILING DATE: 2003-05-23

PRIOR PLILING DATE: 2003-04-16

PROBLICATION NUMBER: US 10/417,012

PRIOR PLILING DATE: 2003-04-16

PRIOR PLILING DATE: 2005-14:36:14

PRIOR PLILING DATE: 2002-05-23

PRIOR PLILING DATE: 2005-14:36:14

PRIOR PLILING DATE: 2002-05-23

PRIOR PLILING DATE: 2005-14:36:14

PRIOR PLILING DATE: 2002-05-23

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